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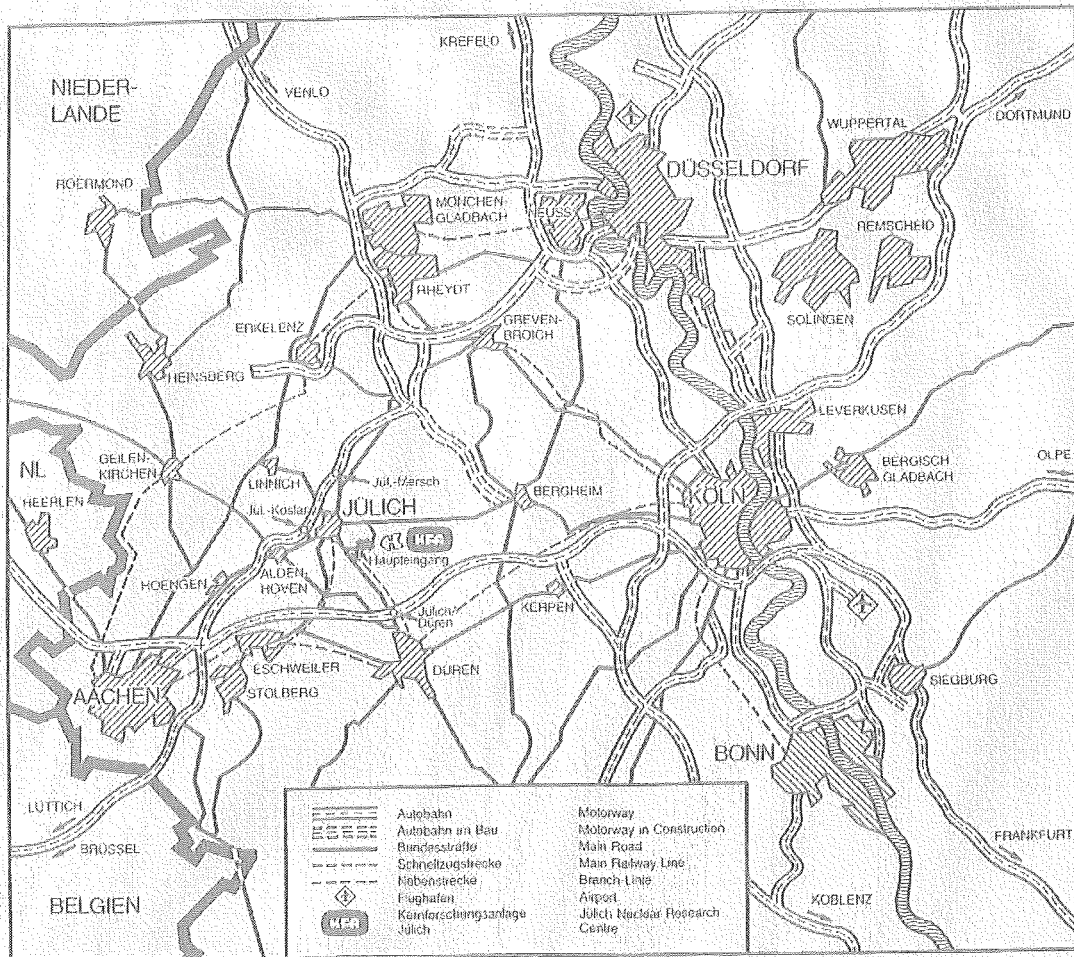
Institut für Kernphysik

**BIG KARL MAGNET SPECTROMETER**  
**Operating System**  
**User's manual**

by

J. Meißburger

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**BIG KARL MAGNET SPECTROMETER**  
**Operating System**  
**User's manual**

by

J. Meißburger

## I n t r o d u c t i o n

The high resolution magnet spectrometer BIG KARL is in operation since early 1979 at the nuclear research institute IKP of the KFA Jülich. A large number of experiments of different type and varying complexity has been carried out since then mostly in close collaboration between physicists of the IKP magnet spectrometer staff and external visiting groups.

The large number of instrumentation parameters and the sophisticated ionoptical optimization procedures clearly demand for computer control and operating support especially if one considers the large distances between spectrometer area, cyclotron- and experimental control rooms.

On the other hand the rather complex detector and data acquisition systems require a computer with sufficient computing power and high speed peripheral I/O to provide on-line control of the experiment, real-time data handling capabilities and a significant amount of general support for preparing and performing an experiment. Finally the rough experimental data have to be processed to deliver the concentrated physical results ready for publication.

The Big Karl operating system comprises two computers, one small DEC PDP-11/40 system for control and a VAX-11/780 (the only 32-bit "supermini" available at that time) for data acquisition and analysis.

### 1) Spectrometer Control

Computer control is handled by a PDP-11/40 computer running RSX-11M which controls all spectrometer components such as movable slits, field measurement probes, magnet currents or general purpose multiplexed DVM's. In addition this computer does automatic background logging of all control activities in order to generate status reports or to display and analyze measured magnetic field data for ionoptical optimization. The main tasks to do these jobs are:

CYCLE	Spectrometer control main task
-------	--------------------------------

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Author: J.Meissburger

STATUS	Spectrometer status report generation
CYSTOP	Unsolicited operating system reset (emergency)
RESET	Control system soft reset
ZEROPS	Control system database initialization
FUMMEL	Magnetic field measurement task
FUPLOT	Interactive field measurement plotting and analyzing utility
SAVBATCH	Procedure to save log files
SAVDATA	Procedure to save data files

CAMAC interface testing is done automatically at system startup. In addition there is a set of tasks available for testing single or complex components of the control system. (Examples are TTCYC wich tests the CYCLE global section and backup file consistency, TTSPRE which tests the iterative direct field setting of the main quadrupoles, or TTHT which tests the control of the 24 Ht power supplies)

More detailed information on the system is available on-line using the commands:

```
MCR> HELP      System help      and
MCR> HELP      Applications help
```

## 2) Data Acquisition and Analysis

A special hardware system MEMPHIS with fast hardware preprocessing and windowing capabilities has been developed by the KFA's central laboratory of electronics ZEL. It interfaces up to 16 high resolution ADC's, TDS's or pattern units via CAMAC to a VAX unibus. The software utilities

MEMPHIS	MEMPHIS hardware configuration
MPSTATUS	and hardware status logging

# BIB KARL magnet spectrometer operating system HELP LIBRARY

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are available for user-friendly and self-documenting definition and downloading of hardware configurations to the microprocessor-controlled MEMPHIS system.

The main data acquisition tasks (ACQUIRE) are:

ACON	ACQUIRE main data taking task
ACPARAM	Pseudo-parameter definition
ACSTATUS	Data taking soft/hardware status

which run on a special dedicated user account BK\_ONLINE. While the MEMPHIS system provides physical and pseudo-ADC's (by the internal hardware processing units) the ACON system allows free user-defined pseudo-parameter definitions (Definition language is Fortran) to handle more complex detector setups such as the Morris-type drift chambers. The interactive utility

MOCAL	Morris chamber calibration
-------	----------------------------

is available for on-line generation of the Morris-chamber calibration tables used in ACON/ACPARAM pseudo-parameter definition.

List mode data taken by ACON may be reanalyzed by the off-line sorting utilities

PLSORT	General purpose off-line sorting
ACOFF	Off-line version of ACQUIRE

where PLSORT offers various features such as batch mode sorting, one-dimensional and scatter plot displays, complex gate definitions and ACPARAM-compatible pseudo-parameters. In order to document or further process the sorted spectra the utilities

ACPLOT	Line plotting of spectra
ACSPECT	Spectrum fitting and analysis
BKMERGE	Spectrum manipulation and display



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are available to accept true physical or Monte-Carlo-generated spectra and produce any kind of combined or user-manipulated pseudo-spectra, enhanced screen or hardcopy displays and extracted physical summary data (fits) which are input to the physical data analyzing system

BKDAP	Big Karl data analysis programs, Cross sections, angular distributions and DWUCK
-------	--

### 3) Spectrometer operating support

In parallel to the on-line control and data acquisition tasks a set of utilities for experimental setup and ionoptical spectrometer tuning is available:

BKARLO	Particle Energy loss trough layered layered detector setups
BRHO	General purpose kinematics with muptiple targets and backing
BKSEARCH	Calibration reactions crossover search
MAGSET	Semi-empirical magnetic field setting
MATRIX	Experimental raytracing and spectrometer optimization utility
KIN1	Spectrometer K-value tuning

### 4) General User Support

In addition to these more specific tasks a set of commonly used programs for spectrometer performance tuning and general purpose data analysis are installed:

TRANSPORT	CERN/SLAC TRANSPORT ionoptical code
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TURTLE	CERN TURTLE-decay
MAD	CERN MAD ionoptical design program
STRAGGLE	Energy straggling (Vavilov distribution)
MINUIT and MINFIT	CERN general purpose fitting utility with enhanced user support (MINFIT
FOWL	CERN FOWL phase space calculation
IGM	Big Karl Interactive Graphics Monitor
CTR	COSYTRIEVE database manager (Publication index or CAMAC module database)

For Fortran programmers a collection of object libraries is found on a dedicated directory BKLIB: which comprises

CERN	The CERN library (VMS version)
GMAGII (AGII)	Tektronix Advanced Graphing II with enhanced
GMA10 (PLOT10)	PLOT10 terminal control system (TCS)
CALCOMP and CALTRON	CALCOMP line plotter library and Calcomp emulator for Printronix graphics printer,
BKUTY	Big Karl utility subroutines

which offer a set of BK TT.... routines for general system support particularly for user-transparent terminal configuration in graphic applications and the RD... routines for free-style terminal I/O with intrinsic logging, batch, journal/recovery and error-tolerant modes. The RD-routines are thoroughly used by the Big Karl utilities for standard command parsing.

The by far most important support on the system is provided by the extensive Big Karl HELP library which allows keyword-oriented access by either the

\$ BKHelp      Help command at DCL level

or by a HELP-command at program command level and which documents the entire



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Big Karl operating system down to command parameter and library subroutine source header level. Details on the BKHelp command and how to produce hardcopies are found with

\$ BKHelp BKHELP...

The basic operating principles of the spectrometer Big Karl and its focal plane detectors are summarized in the following three publications which are mandatory for anyone wishing to do a Big Karl experiment:

- 1) Der Magnetspektrograph BIG KARL  
und die Untersuchung des Übergangskernes  $^{103}\text{Ru}$   
mit (d,p)- und (p,d)-Reaktionen  
W.Hürlimann,  
Thesis, KFA internal report, Jül-Spez-104, April 1981
- 2) The QDDQ Magnet Spectrometer "BIG KARL"  
Nucl. Instr. and Meth. 214 (1983) 281-303
- 3) The new focal plane detector  
for the magnet spectrometer Big Karl  
Nucl. Instr. and Meth. A242 (1985) 95-102

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP HELP (14 July 1986):

HELP

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\* BIG KARL magnet spectrometer operating system \*

\*\*\*\*\*

H E L P     L I B R A R Y

This library provides Big Karl users with the necessary information  
to operate the magnet spectrometer control and data acquisition system.

For more details on the BKHELP command use "\$ BKHelp BKHELP" !

Author: J.Meissburger, KFA-IKP,

Last library update: 14 July 1986

Additional information available:

ACPLOT	ACQUIRE	BKARLO	BKFIT	BKHelp	BKLIB	BRHO
CERN	COSYTRIEVE	CTR	CYCLE	FUMMEL	FUPLOT	Graphics
HELP	IGM	MATRIX	MEMPHIS	MINIMI	MINUIT	MOCAL
NEWS	PATCHY	PLSORT	PRINTERS	RDSHOW	RSX11M	RUNOFF
System_layout		Telephone	Terminals	Utilities		

Author: J.Meissburger

BKHELP

\*\*\*\*\*

H E L P      L I B R A R Y

This library provides Big Karl users with the information needed to operate the magnet spectrometer computer control and data taking systems. It includes:

To keep track of recent developments it is strongly recommended to use:     \$ NEWS       or       \$ BKHelp NEWS  
at least once before starting a new experiment !

To access the help information type:

```
$ BKHlp      key1, key2, .....
```

where key1, key2 are the search keys listed under "further information available:"

or

```
$ BKHelp/output=BKHELP.LIS      key1, key2, .....      and
$ Print/delete BKHELP.LIS
```

to get a hardcopy of the BKHELP information.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Remember: Despite all documentation a basic understanding  
of magnet spectrometers and computers is highly  
desirable for performing a Big Karl experiment !

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BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP ACPLLOT (14 July 1986):

ACPLLOT

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\*\*\* ACPLLOT \*\*\*

\*\*\*\*\*

Author: B.Brinkmoeller

The Program ACPLLOT may be used to make a histogram plot of spectra stored in \*.SIN - files.

The output device is either the CALCOMP PLOTTER or the PRINTRONIX LINE PRINTER/PLOTTER.

valid commands are :

READ : to read data from a \*.SIN - file  
SET : to set new paramters for the plot  
STATUS : to see the present parameters and the data stored  
HELP : to get informations about the program  
PLOT : to plot arrays of the stored \*.SIN - files  
EXIT : to exit to DCL level  
PAGE : to start following plot on new page

ACPLLOT

READ

the READ command is used for data transfer to the program

You have to give the name of the data file to read.

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Author: J.Meissburger

It will prompt you for the array numbers you wish to transfer.  
If you do not give any array number the first sixteen arrays of  
the file will be transfered to the program.

The maximum number of arrays that can be stored at one time is sixteen.

## ACPLOT

### SET

with the SET command you can change the following parameters :

XMIN	the minimum channel number for the plot
XMAX	the maximum channel number for the plot
PEN	the number of the pen used for plotting
LENGTH	the length of the plot
HEIGHT	the height of the plot
YMIN	the minimum y - value displayed
YMAX	the maximum y - value displayed
COM	compress factor

The program will prompt you for new values of each parameter

If you press <RETURN> the parameter value is not changed

To suppress the prompting you may specify the parameters you want  
to change in the form: "SET LEN 30 XMA 1000"

Only the parameter LENGTH and XMAX are changed and the program returns  
to command level

## ACPLOT

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Author: J.Meissburger

SET

XMIN

If you enter the parameter "XMI"  
you may change the minimum channel number for the plot

You are not allowed to give a minimum x - value larger than  
the maximum x - value unless XMAX is not equal to zero

If XMIN is equal to zero the program sets the minimum x - value  
equal to one

ACPLOT

SET

XMAX

If you enter the parameter "XMA"  
you may change the maximum channel number of the plot

You are not allowed to set XMAX lower than XMIN

If XMAX is equal to zero the program sets the maximum x -value  
equal to the size of the array to be plotted

ACPLOT



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SET

PEN

If you enter the parameter "PEN"  
you may change the number of the pen used for plotting the output

Pen numbers in the range from 1 to 4 are allowed  
the pen number does not have any influence if the output device is the  
line printer

ACPLOT

SET

LENGTH

If you enter the parameter "len"  
you may change the length of the plot

There is no restriction on the maximum length

ACPLOT

SET

HEIGHT

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Author: J.Meissburger

If you enter the parameter "HEI"  
you may change the height of the plot

the height of the plot is limited by the paper - width on  
the output device that you choose

ACPLOT

SET

YMIN

If you enter the parameter "YMI"  
you may change the minimum y-value for the output

You are not allowed to give a minimum y - value larger or equal to  
the maximum y - value unless XMAX is equal to zero

If YMIN is equal to zero the program will set the minimum y - value  
to zero or the minimum negative value in the array to be plotted

ACPLOT

SET

YMAX

If you enter the parameter "YMA"  
you may change the maximum y - value for the output

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Author: J.Meissburger

You are not allowed to give a maximum y - value lower than or equal to the minimum y - value unless you set it to zero

If YMAX is equal to zero the program sets the maximum y - value to zero or the maximum positiv value in the array to be plotted

ACPLOT

SET

COMPRESS

If you enter the parameter "COM"  
you may change the compress factor of the spectra

The maximum compress factor is 99

ACPLOT

SET

DEFAULT

If you enter "DEF"  
the parameters of the plot are reset to defaults that were  
found suitable for the output device you choose

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Author: J.Meissburger

ACPLOT

SET

RETURN

If you enter "RET"  
you return to the command level

Everything you enter after the "RET" is interpreted as an ACPLOT command

"RET" is usefull to supress prompting for parameters you do  
not want to change

It has to be used if you want to enter ACPLOT commands after the "SET"  
command on the same textline.

ACPLOT

EXIT

Exit to DCL - level

If the output - device is the PRINTRONIX LINE PRINTER PLOTTER this command  
submits the output to the printer queue

ACPLOT

STATUS

## BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

With the STATUS command you can display the parameters specifying the output and information about the data stored

### ACPLOT

#### PLOT

With the PLOT command you can produce histogram plots of the spectra on the output device you choose

The program prompts you for the numbers of the arrays you want to plot. If you do not give any array number the program will plot all arrays stored

To change the parameters of the plot like LENGTH and HEIGHT and minimum and maximum x - and y - values, use the SET - command before using the plot command.

### ACPLOT

#### PAGE

The PAGE command enforces output of the following plot on the next page.

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Author: J.Meissburger

BKHELP ACQUIRE (14 July 1986):

ACQUIRE

\*\*\*\*\*

\* A C Q U I R E \*

\*\*\*\*\*

BIG KARL data acquisition utility

Authors: D.Kouzes, R.Weber, H.Stoff, J.Meissburger

The utility is invoked by two DCL-level commands:

\$ ACON	for on-line data taking with MEMPHIS
\$ ACOFF	for off-line default sort and display

It is supported by additional procedures AC... to facilitate tape handling and file management. These procedures are strongly recommended to be used in order to guarantee unique file and tape formats!

The ONLINE utility ACON can o n l y be run by the turnkey-user

BK\_ONLINE (UIC = [100,100])

and produces its data files on directory USERON. These files have to be saved by the user to either magtape or any of the private directories.

ACQUIRE

EXAMPLE

LOGIN

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Author: J.Meissburger

\*\*\*\*\*

To start data acquisition with Big Karl's VAX/MEMPHIS log in as:

username: BK\_ONLINE

password: ..... (ask BK-Manager)

This puts you onto the on-line directory USERON and in addition displays some information on required data files:

MPSxxx = MEMPHIS configuration batch file

MOCxxx = Latest MOCAL calibration file

and a Fortran source file to define BK\_USER\_INI for ACQUIRE parameter setup initialization. Keep a hardcopy of these filenames !

Now perform a few checks before starting:

```
$ SHOW QUOTA    check for free disk space, >10000 blocks !
$ SHOW DEVICE   check for online condition and error counts on:
                DRA1:   Userdisk
                DUA1:   Data disk
                CAxx:   Camac
                GMA0:   GMA display
```

+-----

SETUP MEMPHIS

\*\*\*\*\*

Logon as BK\_ONLINE on the BK-VAX and type:

\$ MEMPHIS

MEM> SET COIncidence 1 SET COIncidence 2 [SET NOCoincidence xx]

MEM> EXIT



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.....

MPSLINK> EXIT

\$ MPStatus        Check physical ADC range switches,  
                  MEMPHIS ADC configuration and coincidence !

If necessary perform a hard reset by pressing the RESET-button on the  
MEMPHIS microprocessor unit (rightmost module, use pencil !) and  
reload configuration with MPS.

+-----

MOUNT A NEW MAGTAPE

\*\*\*\*\*

Magtapes to be used by ACQUIRE must be mounted b e f o r e  
invoking ACON ! Type:

\$ ACINI

....            Default label is the volume-id  
                 BKxxxx (xxxx = green sticker on tape)  
                 Default density (<RET>) = 1600 bpi

This initializes (empties and lables) the tape and mounts it for use  
by ACQUIRE.

+-----

START ACQUIRE

\*\*\*\*\*

Now you are ready for taking data:

\$ ACONline

.... <RET>        No new realtime process (user parameters)

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```
<RET>      Magtape already mounted
GMAO:      plot terminal (<RET> = your terminal)
<RET>      511 events per buffer (obligatory with tape !)

ACON> RE,D
.... <RET>  read last MAIL file to restore spectra setup
ACON> TAPE  switch on list mode data taking on magtape
```

+-----

PRODUCTION LOOP

\*\*\*\*\*

\*\*\*\*\* Data taking loop for production starts here \*\*\*\*\*

```
ACON> ZA,A      Clear old spectra
ACON> AC        Start datataking

....           Please write a short comment to your list mode
                file containing typically at least:
                Target type
                Reaction type
                Lab angle
                Special comments

<RET>
ACON>          <<<<<<<< data taking is now active ! >>>>>>>
```

If you need to stop data taking for any reason (to check electronics, system overload by high intensity etc.) you should use the BIG SWITCH on the NIM crate. This virtually holds real time without changing any hard- or software condition.

For longer STOPS you better stop data taking and close the data file:

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```
ACON> ST          stop data taking
ACON> EF          Close your list mode file (End File)
ACON> DS          Display scalers >>>> L O G B O O K !
                  >>>> Make a hardcopy of the borer scalers
                  and make notes in the L O G B O O K !
```

```
ACON> DU,D        write your sorted spectra to disk
....             comment see above
```

\*\*\*\*\* Data taking loop for production ends here \*\*\*\*\*

+-----

PRODUCTION END

\*\*\*\*\*

Close and dismount the magtape

If you are near the end of your tape or if you want to continue  
on a new tape:

```
ACON> NT          Disable list mode tape
ACON> DU          Dump mail file
....             Comment: FOR PARAMETER SETTING
ACON> EXIT
```

Do you want to keep the on-line process running (Y/N): NO

~~\$ ACDISmount~~ Dismount/nounload your tape

Take the write-ring off and put the tape back in the cupboard !  
Get the next free tape (see above)

In order to save a l l data files important to your experiment on  
a backup tape it is strongly recommended to use:

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## \$ ACSAVE

after finishing your run. ACSAVE not only saves the on-line spectra but also the software and hardware configuration files as well as any parameter definitions or calibration tables.

+-----

## TAPE REQUESTS

\*\*\*\*\*

In case you run into troubles with a magtape service (e.g. unexpected End Of Tape) you may login as:

username: OPERATOR

(no password)

The system displays a list of outstanding tape service operator requests. Cancel your request by typing in the corresponding REQUEST-ID number  
b e f o r e    you exit ACQUIRE !!!!!!!!!!!!!

ACQUIRE

COMMANDS

ACQUIRE

COMMANDS

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SUMMARY

Alphabetical Summary of ACQUIRE Commands

03-OCT-81

-----

Assignments:     DUMP\$DEVICE = default disk  
                  EVENT\$DEVICE= default disk  
                  or    EVENT\$DEVICE= MAGTAPE if tape is mounted

type:    ..ACQ>HELP ERRORS            for error messages

type:    ..ACQ>HELP xxxx            where xxxx is any legal command

@filename	Start command file 'FILENAME.BAT'
2D	Create a two dimensional array
ACQ	Start data taking from MEMPHIS
ADC	Set ADC associated with array
ARRAY	Define one dimensional arrays
BELL	Ring bell
CLEAR	Clear Graphic Scope screen (after SCOPE)
CONDITION	Set side conditions on storing data
DARRAY,n	Delete array N
DSCALERS	Read and display current scaler values
DUMP	Dump parameters and arrays to DUMP\$DEVICE
DUMP,D	Dump parameters and arrays to DUMP\$DEVICE
DUMP,E	Dump parameters and arrays to EVENT\$DEVICE
EFILE	Write an END OF FILE on magnetic tape
EXIT	Exit the program
GAINS	Set ADC conversion gain ( only for 2D arrays )
LIVE	Show the live data coming into an array
MSCALER	Manipulate Preset Scaler
NOTAPE	disable list mode on magtape
OFFSET	Set array zero offsets
PARAMETERS	List the actual data taking parameters

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PARAMETERS,P	Print the actual data taking parameters on line printer
PLOT,n	Printer plot of 1-D array N
PRINT,n	Print array N on lineprinter.
READ	Read data arrays or list mode data from DUMP\$DEVICE
READ,D	Read data arrays or list mode data from DUMP\$DEVICE
READ,E	Read data arrays or list mode data from EVENT\$DEVICE
READ,O	Read data arrays or list mode data from other directory
SCOPE	Display array on Graphic terminal
SCOPE,n	Display array N on Graphic terminal ( prev. scale )
SINGLES,n	Set Singles mode of data taking ( default )
STOP	Stop data taking from MEMPHIS
SUM	Sum data array shown on scope within limits (Cursor)
SUM,A	same, but enter channel numbers from keyboard
TAPE	Enable list mode data taking on magtape
ZAP,n	Set array N to zero
ZAP,ALL	Set all arrays to zero

## ACQUIRE

## COMMANDS

## ERRORS

If the user enters an invalid command, the program outputs a message

ERR?xx, where xx is:

BT	Bad Tape
SP	Write Protected
PE	Parity Error
CC	Invalid control command

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Author: J.Meissburger

NC	No more core available
AB	User request aborted
ET	End of magnetic tape
NP	No parameters have been entered
MT	Magnetic tape error
NS	No SAVE command given
TP	Too many parameters
LP	Printer error or tape at load point
DK	Disk error
EV	Event mode task error
FL	Digital input flag error
OP	Error on OPEN attempt

ACQUIRE

COMMANDS

@

@filename        Start command file 'FILENAME.BAT'

-----  
command file must be created with: \$ EDI filename.BAT  
example of a command file:

SINGELS,1

ARRAY

1

1

2048

-

-

2D



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Author: J.Meissburger

2h

1,128,0,32

2,128,0,32

—

—

PARAMETERS

ZAP,ALL

ACQUIRE

COMMANDS

2D

2D                      Create a two dimensional array

---

ACQ>2D

ARRAY #? 5\_ \

X ADC, SIZE, OFFSET, GAIN:7,128,0,16

Y ADC, SIZE, OFFSET, GAIN:8,32,0,32

CONDITION ADC #? \_ \

ARRAY #? \_\_ \

ACQ>

ACQUIRE

COMMANDS

ACQ

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Author: J.Meissburger

ACQ                    Start data taking from MEMPHIS

---

ACQUIRE

COMMANDS

ADC

ADC                    Set ADC associated with array

---

ACQ>ADC

ARRAY #? 2\_\

ADC #? 8\_\

ARRAY #? \_\_\

ACQ>

ACQUIRE

COMMANDS

ARRAY

ARRAY                    Define one dimensional arrays

---

ACQ>ARRAY

ARRAY #? 3\_\

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

ADC #? 3\_\

SIZE? 512\_\

CONDITION ADC #(0 TO CLEAR) \_\

ARRAY #? 4\_\

ADC #? 4\_\

SIZE? 512\_\

CONDITION ADC #(0 TO CLEAR) \_\

ARRAY #? \_\_\

ACQ>

## ACQUIRE

### COMMANDS

#### BELL

BELL                      Ring bell

-----

## ACQUIRE

### COMMANDS

#### CLEAR

CLEAR                      Clear Graphic Scope screen

-----

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

ACQUIRE

COMMANDS

CONDITION

CONDITION           Set side conditions on storing data

-----

ACQ>CONDITION

ARRAY #? 1\_\

CONDITION ADC #(0 TO CLEAR) 3\_\

LOWER LIMIT(4K SCALE), OPTIONAL MULTIPLIER

200

UPPER LIMIT(4K SCALE), OPTIONAL MULTIPLIER

500

CONDITION ADC #(0 TO CLEAR) 4\_\

LOWER LIMIT(4K SCALE), OPTIONAL MULTIPLIER

50

UPPER LIMIT(4K SCALE), OPTIONAL MULTIPLIER

100

CONDITION ADC #(0 TO CLEAR) \_\

ARRAY #? \_\_\

ACQ>

ACQUIRE

COMMANDS

DARRAY

DARRAY,N

Delete array N

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

---

ACQ>DARRAY  
ARRAY #? 2\_\n  
ACQ>

ACQUIRE

COMMANDS

DSCALERS

DSCALERS            Read and display current scaler values

---

ACQUIRE

COMMANDS

DUMP

DUMP                Dump parameters and arrays

---

DUMP                Dump parameters and arrays to DUMP\$DEVICE  
file is MAIL.SIN for default mailbox file

DUMP,D              Dump parameters and arrays to DUMP\$DEVICE  
filenames are automatically updated to ACxxxx.SIN

DUMP,E              Dump parameters and arrays to EVENT\$DEVICE  
filenames are automatically updated to ACxxxx.SIN

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

ACQUIRE

COMMANDS

EFILE

EFILE                    Write an END OF FILE on magnetic tape

---

close list mode file on magnetic tape

ACQUIRE

COMMANDS

EXIT

EXIT                    Exit the programm

---

NOTE:    EXIT does    n o t    stop the subprocess created by  
         ONLINE Acquire. To stop the subprocess, use \$ ACSTOP !!!!

ACQUIRE

COMMANDS

GAINS

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

GAINS                    Set ADC conversion gain ( default by MEMPHIS )

---

ACQ>GAINS

ARRAY #? 1\_ \

GAIN? 1\_\_ \

ARRAY #? \_\_ \

ACQ>

ACQUIRE

COMMANDS

LIVE

LIVE                    Show the live data coming into an array

---

ACQUIRE

COMMANDS

MSCALER

MSCALER                Manipulate Preset Scaler

---

ACQ>MSCALER

VALUE? 1000\_\_ \

ACQ>



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

ACQUIRE

COMMANDS

NOTAPE

NOTAPE            disable list mode on magtape

---

ACQUIRE

COMMANDS

OFFSET

OFFSET            Set array zero offsets

---

ACQUIRE

COMMANDS

PARAMETERS

PARAMETERS        Print the data taking parameters

---

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

PARAMETERS,P      Print the data taking parameters on lineprinter

PARAMETERS        Type the data taking parameters on terminal

ACQUIRE

COMMANDS

PLOT

PLOT,n            Printer plot of 1-D array N

---

ACQUIRE

COMMANDS

PRINT

PRINT,n           Print array N on lineprinter

---

ACQUIRE

COMMANDS

READ

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

READ                    Read data arrays or list mode data

---

READ                    Read data arrays or list mode data from DUMP\$DEVICE  
                         <RET> for filename reads the default mailbox file

READ,D                  Read data arrays or list mode data from DUMP\$DEVICE  
                         <RET> for filename reads the default mailbox file

READ,O                  Read data arrays or list mode data from DUMP\$DEVICE,  
                         but from OFFLINE directory when running ONACQ  
                         and from ONLINE directory when running OFFACQ  
                         <RET> for filename read the default mailbox file

READ,E                  Read data arrays or list mode data from EVENT\$DEVICE

ACQUIRE

COMMANDS

SCOPE

SCOPE                   Display array on Graphic terminal

---

SCOPE                   Display array on Graphic terminal

SCOPE,n                Display array N on Graphic terminal ( prev. scale )

ACQUIRE

COMMANDS

SINGLES

BIB KARL magnet spectrometer operating system HELP LIBRARY  
Author: J.Meissburger

SINGLES,n            Set Singles mode of data taking ( default )

---

ACQ>SINGLES,6

ACQ>

ACQUIRE

COMMANDS

STOP

STOP                Stop data taking from MEMPHIS

---

ACQUIRE

COMMANDS

SUM

SUM                Sum data array shown on scope

---

SUM                Sum data array shown on scope within limits (Cursor)  
SUM,A              same, but enter channel numbers from keyboard

The following subcommands are accepted by SUM:

- 0 = EXIT back to main command level
- 1 = MARK channel position with cursor

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

2 = REPLOT  
3 = SCALE up  
4 = SCALE down  
5 = SUM channels and print LSQ-statistics  
6 = Ask for CHANNELS  
7 = MARK and subtrakt background  
8 = Perform Gauss FIT  
  
K = KILL (delete) Matrix file ACMTX.TMP  
R = SUM and add record to ACMTX.TMP  
B = BLANK line added to ACMTX.TMP

ACQUIRE

COMMANDS

TAPE

TAPE                    Enable list mode data taking on magtape  
-----

ACQUIRE

COMMANDS

ZAP

ZAP                    Set array to zero

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

-----

ZAP,n	Set array N to zero
ZAP,ALL	Set all arrays to zero

ACQUIRE

PROCEDURES

ACQUIRE COMMAND PROCEDURES: 21-JUL-83

=====

>>>>> These procedures are available o n l y if logged on  
>>>>> as a Big Karl group user BK\_....

ACQUIRE

PROCEDURES

ONSET

\$ ONSet --> Set default to online directory USERON

ACQUIRE

PROCEDURES

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

ACONline

\$ ACONline --> Start ONLINE Acquire program on directory USERON

ACQUIRE

PROCEDURES

ACSTOP

\$ ACSTop --> Stop ONLINE Acquire subprocess  
>>> Exiting the ACQUIRE program does N O T  
>>> stop the related ONLINE subprocess !!!!!

ACQUIRE

PROCEDURES

ACOFFline

\$ ACOFf --> Start OFFLINE Acquire program

ACQUIRE

PROCEDURES

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

ACPARAM

\$ ACParam --> Define ONLINE user parameters (pseudoparameters)

ACQUIRE

PROCEDURES

ACSTATUS

\$ ACStatus --> Show the actual ONLINE parameter and MEMPHIS status

ACQUIRE

PROCEDURES

ACSPECT

\$ ACSpect --> Start OFFLINE fit program SPECT

ACQUIRE

PROCEDURES

ACPLOT



# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
$ ACPlot    -->   Plot spectra (*.SIN) on CALCOMP-1012 plotter
                  or on Printronix dot matrix printer
>>>   Before plotting on CALCOMP plotter
>>>   check plotter status first (line switch) !!!
>>>   Then press:      RESET (rear panel, light flickering)
>>>                      INDEX (>2 seconds!)
>>>                      MODE (lighth stays stable)
```

ACQUIRE

PROCEDURES

ACDUMP

```
$ ACDump    -->   Dump data files in ACQUIRE format
```

ACQUIRE

PROCEDURES

ACINIT

```
$ ACInit    -->   Initialise and mount a new magtape
```

ACQUIRE

PROCEDURES

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

ACMOUNT

\$ ACMount --> Mount an initialized and labeled tape

ACQUIRE

PROCEDURES

ACDISMOUNT

\$ ACDismount --> Dismount a magtape

ACQUIRE

PROCEDURES

S15PLOT

\$ S15plot --> Plot old PDP15 spectra sorted on PDP-11/40

ACQUIRE

PROCEDURES

ACSAVE

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

\$ ACSave    -->    Save online data on save device  
                  (give starting date of your experiment)

ACQUIRE

PROCEDURES

REWIND

\$ REWind    -->    Rewind a mounted magtape

ACQUIRE

MEMPHIS

The utility MPS is used to configure the Big Karl data acquisition hardware MEMPHIS for a desired experimental setup. It is invoked by typing \$ MPS at DCL command level when logged on as BK\_ONLINE

To see the actual status of MEMPHIS (ADC configuration and coincidence) type: \$ MPSTATUS

ACQUIRE automatically creates a default batch file MEMPHIS.BAT which always reflects the true hardware configuration of MEMPHIS (not taking into account any user- modification of parameters)

For more information see: \$ BKHelp MEMPHIS...

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP BKARLO (14 July 1986):

BKARLO

\*\*\*\*\*

\* B K A R L O \*

\*\*\*\*\*

Authors: J.Meissburger, R.Korthues

Calculate particle energy loss through the BIGKARL magnet spektrometer and its different detector layers.

New BKARLO data file layout:

The primary input file has extension \*.INP and resides on YOUR private directory.

The \*.MOD module files and \*.DAT data files remain on USERDISK:[USER.BKARLO] and are read from there !

BKARLO

Input\_file

The \*.INP file is the primary data input file to BKARLO. It contains a list of module names and the gas pressure in the module where appropriate (mbar). A modulename is a character string (e.g.: 'MOD1') which must be first on the line. Leading blanks are ignored. If a module contains a gas volume (or vacuum) the gas pressure must be defined on the same line separated by at least one blank character. (Note: Gas pressures cannot

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

be exactly 0 !). Reading of numbers is format-free.

BKARLO

Input\_file

EXAMPLE

modulename:	pressure	comment
'MORRIS'	1000.	Morris chamber
'AIRGAP30'	500.	air gap

BKARLO

Module\_file

A module file (\*.MOD) contains a list of different layers. The total number of layers n of a module is given by the parameter L in the form "L=n". BKARLO expects a list of n lines containing each a character string specifying the layer name, and the thickness of that layer in mm. The layer name must be first on the line besides any optional blanks. The layer thickness is separated by at least one blank on the same line. Reading is format-free.

BKARLO

Author: J.Meissburger

### EXAMPLE

layer	thickness	comment
'MYLAR'	0.005	first layer
'MOGAS1'	2.9	
'AIR'	30	

Data file

In the following block the molecules are described in the form:

```
.          .
.         .                                     one molecule
.        .
.       .
A=am   N=nm    zm ATOM
```

next molecule

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

pn % ...

With:

p = fraction in percent (volume for gases, weight for solids) of one molecule in the layer

a = atomic mass number of contributing atom

n = nuclear charge of contributing atom

z = number of atoms of type A,N present in this molecule

This block is limited by a line containing blanks.

If the program finds a numerical value in one of the following lines before EOF this value will be read format-free and interpreted as layer density.

If the layer represents a plastic scintillator a "1" in the following line tells BKARLO to calculate light output for this layer.

BKARLO

Data\_file

EXAMPLE

FILE: NE110.DAT

Delta-E plastic scintillator

I=-1 number of molecules in the layer, negative for solids

100 % , A= 1 , N=1 , 1104 ATOMS ; HYDROGEN

A=12 , N=6 , 1000 ATOMS ; CARBON

1.03D0 DENSITY

1 SCINTILLATOR LIGHT OUTPUT SWITCH

BKARLO

Theory

The program calculates for each layer the specific compound- or gas-mixture energy loss by adding the weighted contributions of the different atoms in the mixture. The mean density is calculated as for an ideal gas.

Mean ionization potentials are empiric for nuclear charges smaller than 33 and are theoretically calculated for heavier nuclei.

The energy loss of a particle is then integrated as it passes through the layers of the detector setup and loses energy.

For more details ask: J.Meissburger



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP BKFIT (14 July 1986):

BKFIT

\*\*\*\*\*

\* B K F I T \*

\*\*\*\*\*

Authors: J.Meissburger, M.Witteler

The B i g K a r l general purpose analyzing program to  
read , write, merge , modify , analyze and display  
data files written in Big Karl standard format (\*.SIN)

BKFIT

COMMAND\_FORMAT

Commands are executed several in one line from left to  
right and separated by at least one blank according to Big Karl's  
standard free style input:

Three characters minimum are required for command  
recognition except for the special commands:

"!" and ";" which indicate a following comment, and  
"?" which gives a list of valid commands

Numeric input is format-free and is defaulted automati-  
cally to appropriate values or prompted for if required.

## BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Interactive command input format is identical to program level batch mode input and to the journal file output which may be used to reproduce a complete session. In addition all input and output goes to a logging file for later reanalysis.

BKFIT

### PROGRAM\_STRUCTURE

BKFIT is used to make modifications on single spectra or to create new, combined pseudo-spectra. The singles spectra as stored in BIG KARL standard \*.SIN format (e.g. ACQUIRE or BKFIT) are read into the READ buffer by means of the READ command (SHO READ). Part or all of may subsequently SAVED into a SAVE buffer. All actions of merging, modifying or displaying refer to the spectra in the save buffer (SHO SAVE). The save buffer may either be filled from different \*.SIN files or by BKFIT as a result of the manipulations. The REAL\*4 format of the save arrays reduces rounding errors and overflows when manipulating the data.

With the WRITE command all active arrays from the save buffer are written to a file BKF\*\*\*\*.SIN. Here the format is changed back to Integer\*2.

BKFIT

TEXT

Command to switch on the graphics text mode at the position

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

defined by the graphics cursor. Text up to 79 character per line may be entered directly, graphics input mode is terminated by typing <RET> in response to the graphics cursor. Text is displayed in the actual GTINDEX colour.

BKFIT

RECOVER

This command allows to reproduce the last session of BKFIT by reading all commands from the journal file BKFIT.JOU (if saved).

BKFIT

EXIT

Exit of the program to the operating system command level.

BKFIT

READ

Reads single arrays from a data file in Big Karl standard format containing sorted spectra information (\*.SIN).

READ filename [arraynumber,...]

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

The filename with extension has to be given. The file must be written in standard BIG KARL format (i.e. collected by MEMPHIS data acquisition system or \*.SIN files created by PLSORT or BKFIT). These files contain sorted spectra in a special format that c a n n o t be typed on a terminal or lineprinter. (See DUMP)

You may specify the array numbers you want to read separated by commas or single blanks. If ommitted the programm reads the first 16 arrays of the file. Two dimensional arrays are presently skipped and ignored.

BKFIT

READ

Examples

```
READ AC0900.SIN 1,2 5
```

Read Arrays no. 1, 2, and 5 from file 'AC0900.SIN' and store then in the READ buffer. Other arrays in his file will be skipped.

```
READ AC0900.SIN
```

The first 16 arrays from the specified file will be read and stored.

BKFIT

SAVE

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Copies selected arrays out of the READ buffer into the  
SAVE buffer.

SAVE r\_arraynumber,... [to s\_arraynumber,...]

The arraynumbers may be seperated by blanks or by commas.

BKFIT

SAVE

R\_ARRAYNUMBER

R\_ARRAYNUMBER selects a certain array from the READ buffer.  
More than one array can be given in one line. 'ALL' is accepted as  
a valied parameter if all arrays from the READ buffer are to be  
transferred to the SAVE buffer.

BKFIT

SAVE

S\_ARRAYNUMBER

S\_ARRAYNUMBER selects the postion in the SAVE buffer for  
the arrays. S\_ARRAYNUMBER has to be separated by R\_ARRAYNUMBER  
by the word 'TO'. If no S\_ARRAYNUMBER is given the SAVE buffer is  
filled up using all free positions that are available.

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

If a desired position in the SAVE buffer is already occupied the program issues a warning and does not save the array. You have to use the DELETE command to free positions in the SAVE buffer.

## BKFIT

### WRITE

Writes all arrays that are presently in the SAVE buffer to a file in BIG KARL standard form. The Filename is generated using the update facility of BIG KARL in the form of BKFxxxx.SIN.

## BKFIT

### SHOW

Displays a table of contents of the READ, SAVED or INTEGRATIONS buffer and the setup of the graphic output.

Valid forms of the Command are :

SHOW READ => display contents of READ buffer

SHOW SAVE => display contents of the SAVE buffer

SHOW SET => display settings for graphic output

SHOW INT => display integration calculations

## BKFIT

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

DELETE

Deletes array from SAVE buffer.

DEL arraynumber,[....]

Where arraynumber is the number of the array in the SAVE buffer to be deleted. If more than one array number is given they have to be separated by commas or blanks. 'ALL' is accepted as a valid parameter to clear the whole SAVE buffer.

BKFIT

ADD

ADD {array 1} {array 2} {array 3}

Add the content of array 1 to the content of array 2 (channel by channel) and store the result in array 3.

BKFIT

SUBTRACT

SUBL {array 1} {array 2} {array 3}

Subtract the content of array 2 from the content of array 2 (channel by channel) and store the result in array 3.

BKFIT

MULTIPLY

MUL {array 1} {array 2} {array 3}

Multiply the content of array 1 by the content of array 2 (channel by channel) and store the result in array 3.

BKFIT

DIVIDE

DIV {array 1} {array 2} {array 3}

Divide the content of array 1 by the content of array 2 (channel by channel) and store the result in array 3.

BKFIT

AFUNCTION

Add a polynomial function of degree four of the channelnumber to an array (channel by channel).

AFU arraynumber coef\_0,[...,coef\_4]



# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Here 'arraynumber' specifies the array from the SAVE buffer.  
'coef\_0' to 'coef\_4' specify the function. Coeficients which  
are ommitted are defaulted to zerro. The result of the above  
command is:

$$\text{Array(arraynumber)} = \text{Array(arraynumber)} +$$
$$(\text{coef\_0} + \text{coef\_1} * \text{channel\_nr.} + \dots + \text{coef\_4} * \text{channel\_nr}^{**4})$$

BKFIT

SFUNCTION

Substract a polinomial function of degree four of the channelnumber  
from an array (channel by channel).

SFU arraynumber coef\_0,[...,coef\_4]

Here 'arraynumber' specifies the array from the SAVE buffer.  
'coef\_0' to 'coef\_4' specify the function. Coeficients which  
are ommitted are defaulted to zerro. The result of the above  
command is:

$$\text{Array(arraynumber)} = \text{Array(arraynumber)} -$$
$$(\text{coef\_0} + \text{coef\_1} * \text{channel\_nr.} + \dots + \text{coef\_4} * \text{channel\_nr}^{**4})$$

BKFIT

MFUNCTION

Multiply a polinomial function of degree four of the channelnumber with an array (channel by channel).

MFU arraynumber coef\_0,[...,coef\_4]

Here 'arraynumber' specifies the array from the SAVE buffer. 'coef\_0' to 'coef\_4' specify the function. Coeficients which are ommited are defaulted to zerro. The result of the above command is:

```
Array(arraynumber) = Array(arraynumber) *
  ( coef_0 + coef_1*channel_nr. + ... + coef_4*channel_nr**4 )
```

## BKFIT

### DFUNCTION

Divided an array by a polinomial function of degree four of the channelnumber (channel by channel).

DFU arraynumber coef\_0,[...,coef\_4]

Here 'arraynumber' specifies the array from the SAVE buffer. 'coef\_0' to 'coef\_4' specify the function. Coeficients which are ommited are defaulted to zerro. The result of the above command is:

```
Array(arraynumber) = Array(arraynumber) /
  ( coef_0 + coef_1*channel_nr. + ... + coef_4*channel_nr**4 )
```

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

## BKFIT

### SET

Changes default settings for graphic output and sets the switch for the user-supplied subroutine BKF\_USER .

Till now the program provides only the possibility to SCOPE an array on a graphics terminal and o produce a local hardcopies where available. For output output on the graphics printer or on Calcomp plotter the program ACPLLOT must be used. To display the actual setting of parameters use SHOW SET.

Command format:

```
SET [plot_number] par_name par_value
```

PLOT\_NUMBER specifies for which of the up to 6 simultaneously displayed arrays the change is to be made. If no plot\_number is given the change is made for all plots.

PAR\_NAME and PAR\_VALUE specify the attributes of the plot that have to be changed. Valid parameters that may be changed are:

```
PLOTNR = number of arrays displayed simultaneously
XMIN   = minimum channel number to be plotted
XMAX   = maximum channel number to be plotted
YMIN   = minimum y - value to be plotted
YMAX   = maximum y - value to be plotted
COMP   = channel compress factor
```

```
MODE   = mode of display (histogram, point plot, panel)
```

```
USER_SWITCH = switch passed to subroutine BKF_USER
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

LINE STYLE = line style (solid, dotted, dashed etc.)

LINE INDEX = line colour index (0..7)

GTINDEX = graphics text colour index (0..7)

BKFIT

SET

PLOTNR

SET PLO number\_of\_plots

This parameter controls the layout for the scope command.  
You may choose between 6 different layouts for the simultaneous display of 1 up to 6 arrays. The location on the screen and size of the individual plots is fixed for each layout.

BKFIT

SET

XMIN

With XMIN you change the minimum channel number for the display.

SET [plot\_number] XMIN channel\_number

PLOT\_NUMBER specifies for which of the up to 6 simultaneously displayed plots the change is to be effective. If no plot\_number is given

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

the change is made for all plots.

CHANNEL\_NUMBER is the new value of the parameter. The program sets a default value of 0.

BKFIT

SET

XMAX

With XMAX you change the maximum channel number for the display.

SET [plot\_number] XMAX channel\_number

PLOT\_NUMBER specifies for which of the up to 6 simultaneously displayed plots the change is to be effective. If no plot\_number is given the change is made for all plots.

CHANNEL\_NUMBER is the new value of the parameter. The program sets a default value of 0. Then the actual maximum channel to be displayed is taken as the length of the array in the SAVE block.

BKFIT

SET

YMIN

With YMIN you change the minimum channel content for the display.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

SET [plot\_number] YMIN channel\_contnet

PLOT\_NUMBER specifies for which of the up to 6 simultaneously displayed plots the change is to be effective. If no plot\_number is given the change is made for all plots.

CHANNEL\_CONTENT is the new value of the parameter. The program sets a default value of 0. If both YMIN and YMAX are set to be 0 the program performs an autoscale to display the full arraycontent.

BKFIT

SET

YMAX

With YMAX you change the maximum channel content for the display.

SET [plot\_number] YMAX channel\_contnet

PLOT\_NUMBER specifies for which of the up to 6 simultaneously displayed plots the change is to be effective. If no plot\_number is given the change is made for all plots.

CHANNEL\_CONTENT is the new value of the parameter. The program sets a default value of 0. If both YMIN and YMAX are set to be 0 the program performs an autoscale to display the full arraycontent.

BKFIT

SET

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

COMPRESS

With COMPRESS you change the compress of an array for display.

SET [plot\_number] COM compress\_factor

PLOT\_NUMBER specifies for which of the up to 6 simultaneously displayed plots the change is to be effective. If no plot\_number is given the change is made for all plots.

COMPRESS\_FACTOR is the number of channels whose content is summed for display. The program sets a default value of 1.

BKFIT

SET

MODE

With MODE you may change the mode of display.

SET [plot\_number] MODE mode\_number

PLOT\_NUMBER specifies for which of the up to 6 simultaneously displayed plots the change is to be effective. If no plot\_number is given the change is made for all plots.

Three modes are available selected by MODE\_NUMBER.

Mode 1 : Histogram plot (Default)

Mode 2 : Bar plot

Mode 3 : Point plot

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKFIT

SET

USER\_SWITCH

Change the value of the switch passed to user written subroutine  
BKF\_USER.

SET USE\*R\_SWITCH

USER\_SWITCH may be used to select different modes of operation of  
BKF\_USER. This way it is possible to perform a number of different tasks  
with one subroutine. The need to compile new versions of BKFIT for each  
of these tasks can be greatly reduced this way.

BKFIT

USER

calls user written subroutine BKF\_USER. If BKF\_USER is not  
given by the user a dummy subroutine is called.

USE\*R {array1} [ {array2} {array3} ]

{array1}, {array2} and {array3} are the numbers of the SAVE arrays  
that are passed to BKF\_USER. For this subroutine the following call  
convention is used.



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
SUBROUTINE BKF_USER ( USER_SWITCH, U_MAX_SIZE, U_SIZE1, U_ARRAY1,  
1                      U_SIZE2, U_ARRAY2, U_SIZE3, U_ARRAY3)
```

C

```
INTEGER*2 USER_SWITCH, U_MAX_SIZE, U_SIZE1, U_SIZE2, U_SIZE3  
REAL*4     U_ARRAY1(*), U_ARRAY2(*), U_ARRAY3(*)
```

The meaning of the variables is the following:

USRE\_SWITCH

input : may be used to select different modes of calculation  
change USER\_SWITCH with the SET USER command  
return: if set to zero prevents any change in SAVE block

U\_MAX\_SIZE

input : maximum allowed channel number for modified array

U\_SIZE1, U\_SIZE2, U\_SIZE3

input : length of arrays taken from SAVE block

U\_ARRAY1, U\_ARRAY2, U\_ARRAY3

input : arrays taken from SAVE block

U\_SIZE1

return: size of modified array

U\_ARRAY1

return: modified array

This Subroutine is only allowed to change the the content and length of {array1}. Before calling BKF\_USER it is checked wether this array is allready used. If so you have to confirme that you want to change it.

On return from BKF\_USER it is check wether U\_SIZE1 is of a reasonable size and if USER\_SWITCH is greater or equal 0. Only if this conditions are fullfilled the changes in the SAVE block are made.

BKFIT

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

## SCOPE

Display arrays on screen of graphic terminals.

SCO\*PE array\_1, [array\_2, ..., array\_6]

Up to six arrays are simultaneously displayed according to the PLOT\_NUMBER set before. The layout for the display is controlled by the SET command.

## BKFIT

## DUMP

DUM\*P array

The DUMP command writes the content of the specified array from the save block to the file BKFDUMP.LIS. This file can be used for listing on a terminal or output on the lineprinter.

It also offers a simple format to transference data to programs which do not offer the possibility to read \*.SIN files.

---

## BKFIT

## COMPRESS

COM\*PRESS array factor

## BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

In contrast to the SET COM\*PRESS comand, which is only relavant for the display of an array, this command actually changes the content of the given array in the Save block. The compress factor may be any real number latrger then 1.

BKFIT

SHIFT

SHI\*FT array channelnumber

With this command it is possible to shift the contence of the given array. A negative value for channelnumber means a shift to the left a positive number a shift to the right. If channelnumber is positiv (negative) the elements on the right (levt) side of the array are lost, the elemnts on the left (right) side are set to 0. Losing array elements by a shift to the right can be prevented by extending the size of the array with the CON\*CATENATE command.

BKFIT

CONCATENATE

CON\*CATENATE arr#1 arr#2 arr#3

A new array (arr#3) is created wich length is the sum of the lengt of arr#1 and arr#2. It contains the elements of arr#1 in the its first part and the elements of arr#2 in the second part.

A special form of this command can be choosen by giving arr#2 = 0.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Then arr#3 is interpreted as channel number and arr#1 is extended by arr#3 channels. These channels are all set to 0. If arr#3 is negative arr#1 is truncated by arr#3 channels.

BKFIT

CLEAR

CLE\*AR used to clear the graphic display

BKFIT

BATCH

BAT\*ch filename

With this command input to the program it can be changed to a batchfile given by filename. The command format of this file is identical to the interactive command format. The last command in the batchfile has to be END\*BATCH to switch command input back to terminal input.

BKFIT

ENDBATCH

END\*BATCH

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Switch input back to terminal input from a command batchfile

BKFIT

INTEGRATE

INT\*EGRATE arraynumber xlow xhigh

With this command input to the program necessary integration-values are calculated between XLOW and XHIGH. Therefore XLOW is the minimum channel content and XHIGH the maximum. All calculationvalues are saved in the INTEGRATION buffer.

In detail you gets the INTEGRALSUM, the PEAKPOSITION, SIGMA, FWHM and the SKEWNESS of the specified array between the xlimits.

BKFIT

OVERLAY

OVE\*RLAY array\_1, [array\_2, ..., array\_6]

Up to six arrays are simultaneously displayed according to the PLOT\_NUMBER set before. The difference between SCOPE and OVERLAY is that SCOPE displays a whole layout with labels, axis, tic marks and the data curve during OVERLAY only draws the actual data curve of the specified array. So, OVERLAY is usefull to displays multiple arrays in the same picture-window in example to compare those arrays.

If you want to overlay for example only one arrays on the second position use a / for the first arrays which should not be

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Author: J.Meissburger

overlayed by an array.

OVE\*RLAY /, [array, ...]

BKFIT

OVERLAY

EXAMPLES

OVERLAY 1 2 3 4

The data curve of the arrays 1,2,3,4 out of the SAVED buffer are displayed on the first four positions.

OVERLAY / 5 // 13

The data curve of array 5 is displayed on the second position during the data curve of array 13 is displayed on the fifth position. The first, third and fourth position are skipped.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP BKLIB (14 July 1986):

BKLIB

\*\*\*\*\*

\* B K L I B \*

\*\*\*\*\*

Big Karl user libraries on BKLIB:

Object libraries:

General purpose:

BKUTY.OLB	terminal I/O, data I/O and system support
BKMAT.OLB	mathematical utilities (MATRIX)

Graphics:

AGII.OLB	Tektronix Advanced Graphing II (I4)
PLOT10.OLB	Tektronix Terminal Control System TCS (I4)
GMAGII.OLB	Advanced Graphing II modified for GMA (NOI4)
GMA10.OLB	Terminal Control System modified for GMA (NOI4) and colour terminal enhancement support
CALCOMP.OLB	Calcomp model 1012 plotter package (terminal line)
CALTRON.OLB	Calcomp model 1012 plotter package emulated on with: CALTRON.DAT character definition file
IGM1DG.OLB,	Interactive Graphics Monitor link library
IGM3DG.OLB,	
IGMGRP.OLB,	
IGMUTY.OLB	

Data analysis:

BKFIT.OLB	BKFIT/BKMERGE link library
-----------	----------------------------

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Author: J.Meissburger

MINUIT.OLB      Minuit with Chisquare FCN library

MINFIT.OLB

MOCAL.OLB      MOCAL Morris chamber calibration link library

PLSORT.OLB      PLSORT sorting utility link library

with : PLCOMM.FOR include file

BKCOMM.FOR include file

ACQINI.OLB      Acquire initialisation library

ACQUIRE.OLB      Acquire main process link library

ACQUIRERT.OLB      Acquire real time subprocess library

with: ACQUIRERT.BIN object module

FOWL.BIN      Object link module for CERN FOWL program

SBMASS.BIN,      Wapstra table object modules

TMASS.BIN

Special device support:

CAMAC.OLB      CAMAC library (ZEL)

\*\*\*\*\*



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKLIB

BKUTY

\*\*\*\*\*

\* Library: B K U T Y \*

\*\*\*\*\*

BIG KARL UTILITY SUBROUTINES

The following modules are available in BKLIB:BKUTY/LIBR

FILUPD            automatic file name creation utility  
LINEPLOT         produce a plot on a non-graphics terminal/printer

BK\_R\_SIN         read singles spectra in ACQUIRE standard format  
BK\_W\_SIN         write singles spectra in ACQUIRE standard format  
BK\_R\_DATA        read data files in ACQUIRE standard format  
BK\_W\_DATA        write data files in ACQUIRE standard format  
BK\_DECODE        decode MEMPHIS events from buffer word

BK\_CPU\_TIME      provide process CPU time used and left  
BK\_WAIT          hibernate process for a given absolute time  
BK\_RAN\_SEED      get a random value for RAN starting seed  
BK\_TT\_MODE       transparent terminal configuration (type unknown)  
BK\_TT\_CONFIG     configure known terminal-type device  
BK\_TT\_TYPE       get device or terminal type from system  
BK\_GET\_PAR       get foreign command parameter list  
BK\_TT\_AST        terminal <Ctrl-Y> AST routine  
BK\_TRNLOG        repetitive translate logical device name

Format free terminal input:

RDSHOW           example of program using free input command parsing  
RDCMD            free input command parser with batch/logging/journal mode  
RDPAR            read a parameter with optional keyword and prompting  
RDPRMT           forced read of a prompted parameter value

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

RD\_HELP acces private HELP library  
RDCOMP similarity factor for error-tolerant terminal input  
RDCONV pack/unpack text arrays and convert lower- to upper case

RDSET set mode for next reading  
RDTEXT read a text string terminated by blank  
RDALL read a text line including blanks  
RDCOMT read a comment line enclosed in apostrophs  
RDDBLE read a REAL\*8 number  
RDFIX read an INTEGER\*4 number  
RDFLT read a REAL\*4 number

\*\*\*\*\*

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKLIB

BKUTY

BK\_CPU\_TIME

```
CC
CC  FILE: BKCPUTIME.FOR      GET CPU TIME VALUES
CC  =====                J.MEISSBURGER, 6-NOV-1985
CC
CC  CALL BK_CPU_TIME ( TIME_USED, TIME_LEFT)
CC
CC  REAL*4 TIME_USED = CPU time used so far by the calling process
CC  REAL*4 TIME_LEFT = CPU time left for the calling process
CC
CC  All times are in seconds. If there is no time limit the
CC  TIME_LEFT = - TIME_USED (e.g. less or equal zero)
CC
CC  *****
```

BKLIB

BKUTY

BK\_RAN\_SEED

```
CC
CC
CC  FILE: BK_RAN_SEED.FOR    Get random seed for Fortran random
CC  =====                number generator
CC                          J.Meissburger , 4-Feb-1986
CC
C
C  SUBROUTINE BK_RAN_SEED (SEED)
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC
C  INTEGER*4      SEED
C
CC
CC Input arguments: none
CC
CC Return arguments:      SEED = integer value to be used
CC                        as starting seed for calls
CC                        to Fortran function RAN
CC
CC Note:
CC      The calling sequence for a Fortran program using
CC      BK_RAN_SEED would be:
CC
CC      main program
CC
CC      INTEGER*4 SEED
CC      .....
CC
CC      CALL BK_RAN_SEED (SEED) ! call once at beginning !
CC      .....
CC
CC      R = RAN(SEED)          ! call any number of times!
CC
CC *****
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKLIB

BKUTY

RDCMD

CC

CC

CC F I L E : RDCMD.FOR COMMAND PARSER

CC ===== J.MEISSBURGER 18-MAY-83

CC

CC

C SUBROUTINE RDCMD(CMDS,LCMDS,NCMDS,PROMPT,

C \* CMD,NCMD,LFLAG)

CC

CC Read a command string prompting with PROMPT and return the first

CC four characters in CMD together with the command number NCMD.

CC (NCMD is to be used for command parsing in a computed GOTO)

CC

CC A question mark '?' as first character on line produces a list

CC of all commands in array CMDS , a semicolon ';' or exclamation

CC point '!' will ignore the entire line as comment string (which

CC nevertheless will be written to batch output if enabled)

CC

CC If an invalid commands is found in program level batch mode,

CC the mode is changed to interactive and remaining commands are

CC skipped.

CC NCMD = 0 is returned when an End Of File is encountered on

CC command input. Control must n o t return to RDCMD after

CC EOF to avoid loops for jobs running in DCL batch mode !

CC

CC LFLAG is a connecting link between the command parser and the

CC called command service routines. It should be returned to RDCMD by

CC all routines as:

CC LFLAG = 2 on very first entry to RDCMD

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```

CC      =      1      on success
CC      =      0      on error in service routine handled
CC                        by the routine itself (message)
CC      =     -1      on unspecific illegal command
CC
CC
CC Input arguments:
CC  INTEGER*4 CMDS(NCMDS)  list of accepted commands
CC  INTEGER*4 LCMDS(NCMDS) length of command to be checked for match
CC  INTEGER*4 NCMDS       total number of commands
CC  INTEGER*4 PROMPT      prompting symbol (3 characters)
CC  INTEGER*4 LFLAG       see above, also return argument !
CC
CC Return arguments:
CC  INTEGER*4 CMD          command symbol
CC  INTEGER*4 NCMD        command number in list
CC  INTEGER*4 LFLAG       see above, also input argument !
CC
CC *****
CC

```

BKLIB

BKUTY

RDPRMT

```

CC
CC
CC  F I L E :   RDPRMT.FOR          PROMPTED READ OF A PARAMETER VALUE
CC  =====
CC                                J.MEISSBURGER 26-MAY-83
CC
C   SUBROUTINE RDPRMT (PROMPT,LENGTH,MODE,MAXBYT,VAL,IFLAG)
CC

```

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```

CC
CC Read a parameter value:
CC The parameter value will be read according to the desired MODE.
CC If a parameter value is not found, or it's type is found to be
CC wrong,
CC or IFLAG is set to -1 on entry to RDPRMT
CC the value will be prompted by the text string PROMPT.
CC
CC INPUT ARGUMENTS:
CC BYTE PROMPT(*)           prompting text
CC INTEGER*4 LENGTH         length of prompt text in bytes
CC INTEGER*4 MODE           type of desired return argument
CC                           = 0      text string
CC                           = 1      I2
CC                           = 2      I4
CC                           = 3      R4
CC                           = 4      R8
CC INTEGER*4 MAXBYT         maximum number of characters returned
CC                           in mode = 0
CC INPUT/RETURN ARGUMENTS:
CC IFLAG                    -1 to force prompting
CC                           >= 0 otherwise
CC
CC RETURN ARGUMENTS:
CC VAL                      =      parameter value (byte, I2, I4, R4 or R8 resp.)
CC INTEGER*4 IFLAG =        -1      on error
CC                           0       success
CC                           >0      number of bytes returned in MODE = 0
CC
CC *****

```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKLIB

BKUTY

RDPAR

```
CC
CC
CC  F I L E :   RDPAR.FOR           READ A PARAMETER KEYWORD AND VALUE
CC  =====
CC                                     J.MEISSBURGER 6-MAR-1985
CC
C   SUBROUTINE RDPAR(PARNAM,LENGTH,MODE,MAXBYT,VAL,IFLAG)
CC
CC
CC  Read a parameter name and value:
CC  The optional parameter name if found will be skipped, the
CC  parameter value will be read according to the desired MODE.
CC  If a parameter value is not given the parameter name is
CC  used to prompt for a value.
CC
CC  INPUT ARGUMENTS:
CC  BYTE PARNAM()   =      optional parameter name
CC  INTEGER*4 LENGTH      length of parameter name in bytes
CC  INTEGER*4 MODE       type of desired return argument
CC                      = 0      text string
CC                      = 1      I2
CC                      = 2      I4
CC                      = 3      R4
CC                      = 4      R8
CC  MAXBYT           = max. number of characters returned in
CC                      mode = 0
CC
CC  RETURN ARGUMENTS:
CC  VAL              =      parameter value (byte, I2, I4, R4 or R8 resp.)
CC  INTEGER*4 IFLAG =      -1      on error
```



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC          0      success
CC          >0     number of bytes returned in MODE = 0
CC
CC *****
CC
```

BKLIB

BKUTY

RDCONV

```
CC
CC
CC  F I L E :   RDCONV.FOR           CONVERT CHARACTER DATA TYPES
CC  =====
CC                                     J.MEISSBURGER 16-AUG-82
CC
C   SUBROUTINE RDCONV(ARR1,NCHAR,MODE,ARR2,IFLAG)
CC
CC  PACK OR UNPACK CHARACTERS STORED IN ARRAY ARR1 INTO ARR2
CC  ACCORDING TO THE DATA TYPE SPECIFIED BY MODE.
CC
CC  IFLAG = 0 IS RETURNED FOR SUCCESS, -1 FOR ERROR.
CC
CC  Input arguments:
CC  ARR1          =      input array (BYTE, I2 or I4 resp.)
CC  INTEGER*4 NCHAR =      number of characters in input array
CC  INTEGER*4 MODE =      0 - lower case to upper case (a .. z --> A .. Z)
CC                                     1 - BYTE to I2 with zero padding
CC                                     2 - BYTE to I4 with zero padding
CC                                     3 - I2 to I4   with zero padding
CC                                     4 - I2 to BYTE
CC                                     5 - I4 to BYTE
CC                                     6 - I4 to I2
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

CC

CC Return arguments:

CC ARR2 = output array (BYTE, I2 or I4 resp.)

CC INTEGER\*4 IFLAG = -1 for ERROR, 0 for success

CC

CC \*\*\*\*\*

CC

BKLIB

BKUTY

RDCOMP

CC

CC

CC

CC F I L E : RDCOMP.FOR SIMULARITY COMPARATOR

CC ===== J.MEISSBURGER 23-OCT-79

CC

CC

C SUBROUTINE RDCOMP(TEXT1,NBYT1,TEXT2,NBYT2,P)

CC

CC

CC COMPARE TWO TEXT STRINGS AND CALCULATE A SIMULARITY FACTOR

CC GIVEN IN PERCENT. THIS MAY BE USED IN COMMAND PARSING PROVIDING

CC A SUFFICIENTLY HIGH DEGREE OF REDUNDANCY ( >4 CHARACTERS )

CC

CC INPUT ARGUMENTS:

CC BYTE TEXT1(),TEXT2() TWO TEXT STRINGS TO COMPARE

CC INTEGER\*4 NBYT1,NBYT2 LENGTH OF TWO STRINGS IN BYTES

CC

CC RETURN ARGUMENTS:

CC REAL\*4 P PERCENTAGE OF SIMULARITY

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

CC

CC \*\*\*\*\*

CC

BKLIB

BKUTY

RD\_HELP

CC

CC

CC F I L E : RDHELP.FOR GET KEYED HELP INFORMATION

CC ===== OUT OF A (USER-) HELP LIBRARY

CC J.MEISSBURGER 14-DEC-82

CC Version: 3-Jul-1986

CC New: System VMS 4.3 and higher: Use Group specific HELP library

CC e.g. BKLIB:BIGKARL.HLB

CC

C SUBROUTINE RD\_HELP (help\_library, level\_1\_key)

CC

CC

CC >>> This routine runs only under VAX-11 VMS

CC

CC This subroutine is to be called after a HELP command at

CC program command level. It opens the HELP library help\_library

CC ('help\_library.HLB') and searches for a level-1 key "level\_1\_key".

CC which usually is identical to the application program name.

CC

CC Input arguments:

CC help\_library character string containing the name of the

CC desired HELP library (\*.HLB)

CC (e.g. help\_library='BKLIB:BIGKARL')

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC level_1_key      character string giving the level-1 key,
CC                  usually the name of the calling appli-
CC                  cation program (e.g. level_1_key='PLSORT')
CC
CC Return arguments:
CC none
CC
CC For additional information see example: $BKHelp RDSHOW...
CC *****
CC
```

BKLIB

BKUTY

RDSET

```
CC
CC
CC
CC  F I L E :   RDSET.FOR           SET READING CONDITIONS
CC  =====
CC                                     AND POINTERS
CC                                     J.MEISSBURGER 15-OCT-79
CC VERS: 7-FEB-1986
CC
C   SUBROUTINE RDSET(IDSET, IVAL, IFLAG)
CC
CC
CC SET AND RETURN OPERATING MODES OF FREE STYLE INPUT ROUTINES RD...
CC
CC INPUT ARGUMENTS:
CC  INTEGER*4 IDSET           MODE SWITCH
CC
CC INPUT/RETURN ARGUMENTS:
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC  INTEGER*4 IVAL, IFLAG  ACCORDING TO FOLLOWING LIST
CC
CC
CC  IDSET = 0          INITIALISE LOGICAL UNITS AND DISABLE BATCH MODES
CC      INPUT PARAMETERS:
CC      IVAL  = LOG. UNIT FOR TERMINAL INPUT  (DEFAULT=5)
CC      IFLAG = LOG. UNIT FOR TERMINAL OUTPUT (DEFAULT=6)
CC
CC  IDSET = 1          SKIP REMAINING CHARACTERS AND PREPARE FOR READING AT THE
CC      BEGINNING OF THE NEXT RECORD (LINE)
CC      RETURN PARAMETERS:
CC      IVAL  = ACTUAL RECORD POINTER OR RECORD JUST READ
CC      IFLAG = -1
CC
CC  IDSET = 2          RETURN ACTUAL TEXT AND RECORD POINTERS
CC      RETURN PARAMETERS:
CC      IVAL  = TEXT POINTER POSITION IN THE CURRENT LINE
CC      IFLAG = RECORD POINTER VALUE
CC
CC  IDSET = 3          SET TEXT AND RECORD POINTERS
CC      INPUT PARAMETERS:
CC      IVAL  = NEW TEXT POINTER VALUE
CC      IFLAG = NEW RECORD POINTER VALUE
CC
CC  IDSET = 4          SWITCH INPUT BATCH MODE
CC      INPUT PARAMETERS:
CC      IVAL  <= 0 SWITCH BATCH MODE OFF
CC      IVAL  > 0 ENABLE BATCH MODE FROM LOG. UNIT IVAL
CC
CC  IDSET = 5          RETURN END OF FILE CONDITION ON INPUT
CC      RETURN PARAMETERS:
CC      IVAL  = ACTUAL RECORD POINTER
CC      IFLAG = 0  IF NO END OF FILE,
CC      IFLAG = -1 IF END OF FILE WAS FOUND ON INPUT
CC
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC IDSET = 6          SWITCH OUTPUT LOGGING MODE (with INPUT LINE ECHO)
CC          INPUT PARAMETERS:
CC          IVAL  <= 0 SWITCH BATCH MODE OFF
CC          IVAL  > 0 ENABLE OUTPUT BATCH MODE TO UNIT IVAL
CC
CC IDSET = 7          RESET TEXT POINTER BY ONE POSITION
CC          PARAMETERS: NONE
CC
CC IDSET = 8          SWITCH LIST KEY (TERMINAL ECHO IN INPUT BATCH MODE)
CC          INPUT PARAMETER:
CC          IVAL >= 0 SWITCH OUTPUT LISTING ON
CC          IVAL  < 0 SWITCH OUTPUT LISTING OFF
CC
CC IDSET = 9          SWITCH COMMAND INPUT JOURNAL MODE (Output)
CC          INPUT PARAMETERS:
CC          IVAL  <= 0 SWITCH JOURNAL MODE OFF
CC          IVAL  > 0 ENABLE JOURNAL MODE ON UNIT IVAL
CC
CC IDSET = 10         SWITCH COMMAND ERROR MODE FOR JOURNAL INPUT
CC          IVAL  <= 0 COMMAND INPUT ABORTED ON ERROR
CC          IVAL  > 0 COMMAND INPUT CONTINUED ON ERROR
CC
CC *****
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKLIB

BKUTY

RDTEXT

```
CC
CC
CC
CC  F I L E   RDTEXT.FOR           READ A TEXT STRING
CC  =====          TERMINATED BY BLANK
CC                               J. MEISSBURGER 15-OCT-79
CC
CC
C      SUBROUTINE RDTEXT (TEXT,NBYTES,MAXBYT,IFLAG)
CC
CC READ AN ASCII STRING OF UP TO 80 CHARACTERS TERMINATED BY BLANK
CC
CC NBYTES IS THE NUMBER OF CHARACTERS READ
CC IF THERE IS NO CHARACTER DIFFERENT FROM BLANK ON THE LINE IFLAG
CC IS SET TO -1
CC INPUT OF UP TO MAXBYT CHARACTERS PRODUCES IFLAG = 0
CC IF THERE ARE MORE THAN MAXBYT CHARACTERS IFLAG IS SET TO +1
CC AND THE REMAINING CHARACTERS ARE SKIPPED
CC
CC THE CHARACTER STRING IS STORED IN THE BYTE ARRAY TEXT
CC
CC INPUT ARGUMENTS:
CC  INTEGER*4 MAXBYT      MAXIMUM NUMBER OF BYTES TO BE STORED
CC
CC RETURN ARGUMENTS:
CC  BYTE TEXT()          BYTE ARRAY CONTAINING TEXT STRING
CC  INTEGER*4 NBYTES      NUMBER OF CHARACTERS READ
CC  INTEGER*4 IFLAG       RETURN FLAG
CC
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

CC \*\*\*\*\*  
CC

BKLIB

BKUTY

RDALL

CC  
CC  
CC  
CC F I L E RDALL.FOR READ A CHARACTER STRING  
CC ===== TO END OF LINE  
CC J. MEISSBURGER 29-NOV-79  
CC  
CC  
C SUBROUTINE RDALL (TEXT,NBYTES,MAXBYT,IFLAG)  
CC  
CC  
CC READ AN ASCII STRING OF UP TO 80 CHARACTERS INCLUDING BLANKS  
CC  
CC NBYTES IS THE NUMBER OF CHARACTERS READ  
CC IF THERE IS NO CHARACTER ON THE LINE, IFLAG  
CC IS SET TO -1  
CC INPUT OF UP TO MAXBYT CHARACTERS PRODUCES IFLAG = 0  
CC IF THERE ARE MORE THAN MAXBYT CHARACTERS IFLAG IS SET TO +1  
CC AND THE REMAINING CHARACTERS ARE SKIPPED  
CC  
CC THE CHARACTER STRING IS STORED IN THE BYTE ARRAY TEXT  
CC  
CC INPUT ARGUMENTS:  
CC INTEGER\*4 MAXBYT  
CC



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC RETURN ARGUMENTS:
CC  BYTE TEXT(80)
CC  INTEGER*4 NBYTES, IFLAG
CC
CC *****
```

BKLIB

BKUTY

RDCOMT

```
CC
CC
CC
CC  F I L E :   RDCOMT.FOR           READ A COMMENT STRING
CC  =====           ENCLOSED IN APOSTROPHS
CC                               J. MEISSBURGER 15-OCT-79
CC
CC
CC
C      SUBROUTINE RDCOMT (COMT,NBYTES,NMAX,IFLAG)
CC
CC
CC  READ A TEXT STRING OF UP TO 79 CHARACTERS ENCLOSED IN APOSTROPHS (')
CC
CC  NBYTES IS THE NUMBER OF CHARACTERS READ AFTER THE FIRST APOSTROP
CC  IF THE FIRST NONBLANK CHARACTER ON THE LINE IS NOT AN APOSTROPH,
CC  IFLAG IS SET TO -1
CC  INPUT OF UP TO NMAX CHARACTERS PRODUCES IFLAG = 0
CC  IF THERE ARE MORE THAN NMAX CHARACTERS IFLAG IS SET TO +1
CC  AND THE REMAINING CHARACTERS ARE SKIPPED UP TO THE NEXT APOSTROPH
CC
CC  THE CHARACTER STRING IS STORED IN THE BYTE ARRAY COMT
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC
CC INPUT ARGUMENTS:
CC  INTEGER*4 NMAX          MAXIMUM NUMBER OF CHARACTERS STORED
CC
CC RETURN ARGUMENTS:
CC  BYTE COMT()            TEXT STRING ENCLOSED IN APOSTROPH
CC  INTEGER*4 NBYTES       NUMBER OF CHARACTERS READ IN
CC  INTEGER*4 IFLAG        STATUS FLAG
CC
CC *****
CC
```

BKLIB

BKUTY

RDFIX

```
CC
CC
CC
CC  F I L E :  RDFIX.FOR          READ AN INTEGER*4
CC  =====
CC                                J.MEISSBURGER 15-OCT-79
CC
CC
C   SUBROUTINE RDFIX (IVAL, ISIG, IFLAG)
CC
CC
CC READ AN INTEGER*4 IN FREE FORMAT GIVING ABSOLUTE VALUE
CC AND SIGN (+ OR - 1) INCLUDING ZEROES
CC
CC IFLAG IS SET TO ZERO IF A VALID INPUT WAS FOUND.
CC IF THE INPUT WAS INVALID, IFLAG IS SET TO -1 AND THE TEXT
CC POINTER IS REPOSITIONED TO THE BEGINNING OF THE TEXT STRING
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC
CC INPUT ARGUMENTS:
CC  NONE
CC
CC RETURN ARGUMENTS:
CC  INTEGER*4 IVAL          ABSOLUTE NUMBER VALUE (TRUNCATED INTEGER)
CC  INTEGER*4 ISIG          SIGN (+ AND - ZERO HAVE DIFFERENT SIGN !)
CC  INTEGER*4 IFLAG          -1 FOR ERROR, 0 FOR SUCCESS
CC
CC *****
CC
```

BKLIB

BKUTY

RDFLT

```
CC
CC
CC
CC  F I L E    RDFLT.FOR          READ A REAL*4 FLOATING POINT
CC  =====          NUMBER
CC                               J.MEISSBURGER 24-OCT-79
CC
CC
C  SUBROUTINE RDFLT (VAL,SIG,IFLAG)
CC
CC
CC READ A REAL*4 FLOATING POINT NUMBER GIVING ABSOLUTE VALUE
CC AND SIGN (+ OR - 1.) INCLUDING ZEROES
CC
CC IFLAG IS SET TO ZERO IF A VALID INPUT WAS FOUND.
CC IF THE INPUT WAS INVALID IFLAG IS SET TO -1 AND THE
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC TEXT POINTER IS REPOSITIONED TO THE BEGINNING OF THE TEXT STRING
CC
CC INPUT ARGUMENTS:
CC  NONE
CC
CC RETURN ARGUMENTS:
CC  REAL*4 VAL          NUMBER ABSOLUTE VALUE
CC  REAL*4 SIG          SIGN (+ AND - ZERO HAVE DIFFERENT SIGN !)
CC  INTEGER*4 IFLAG     -1 FOR ERROR, 0 FOR SUCCESS
CC
CC *****
CC
```

BKLIB

BKUTY

RDDBLE

```
CC
CC
CC
CC  F I L E   RDDBLE.FOR          READ A REAL*8 FLOATING POINT
CC  =====          NUMBER
CC                               J.MEISSBURGER 17-OCT-79
CC
CC
CC
```

```
C  SUBROUTINE RDDBLE (DVAL, DSIG, IFLAG)
```

```
CC
```

```
CC
```

```
CC READ A REAL*8 FLOATING POINT NUMBER GIVING ABOLUTE VALUE
```

```
CC AND SIGN (+ OR - 1DO) INCLUDING ZEROES
```

```
CC
```

```
CC IFLAG IS SET TO ZERO IF A VALID INPUT WAS FOUND.
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC IT THE INPUT WAS INVALID, IFLAG IS SET TO -1 AND THE
CC TEXT POINTER IS REPOSITIONED TO THE BEGINNING OF THE TEXT STRING
CC
CC INPUT ARGUMENTS:
CC  NONE
CC
CC RETURN ARGUMENTS:
CC  REAL*8 DVAL          NUMBER.ABSOLUTE VALUE
CC  REAL*8 DSIG          SIGN (+ AND - ZERO HAVE DIFFERENT SIGNS !)
CC  INTEGER*4 IFLAG      -1 FOR ERROR, 0 FOR SUCCESS
CC
CC *****
```

BKLIB

BKUTY

SBMASS

```
CC
CC  F I L E :   SBMASS.FOR          Return a mass value from
CC          =====                the WAPSTRA 77 mass table
CC                                          I.KATAYAMA 29-JUN-81
CC                                          B.BRINKMOELLER 11.10.83
CC                                          added symbol for neutron
CC
C  SUBROUTINE SBMASS(IA,IZ,AT,ATE,NORAT,SYMBOL)
CC
CC  MASS TABLE IS REPLACED TO NEW WAPSTRA TABLE('77),
CC  WHICH WAS TYPED-IN AND CHECKED BY DR.YAMAZAKI10
CC  AT RCNP,OSAKA UNIVERSITY.
CC
CC  The mass value is returned according to the interger charge and
```

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```

CC mass values. If a negative charge is given a search for the
CC given symbol is made in the table of chemical symbols.
CC
CC INPUT:  IA      integer mass number
CC          IZ      integer nuclear charge number (or SYMBOL)
CC          SYMBOL  chemical symbol of element (or IZ)
CC
CC OUTPUT: AT      nuclear mass less electron masses
CC           ATE    atomic mass including electron
CC          IZ      integer nuclear charge number (or SYMBOL)
CC          SYMBOL  chemical symbol of element (or IZ)
CC FLAG:  NORAT    =1 mass value found
CC           =0    not found
CC
CC Use LINK ...,BKLIB:sbmass.bin,tmass.bin,... to access this routine
CC

```

BKLIB

BKUTY

FILUPD

```

CC
CC F I L E :   FILUPD.FOR           UPDATE SYS-FILE AND RETURN
CC =====
CC                                     NEXT FILENAME
CC                                     J.MEISSBURGER 11-AUG-82
CC
CC SUBROUTINE FILUPD(SYSFIL,NBYT,LSYS,INCR,FILNAM)
CC
CC*****
CC
CC BYTE SYSFIL(NBYT),FILNAM(12)
CC INTEGER LSYS,INCR,NBYT

```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC
CC  input arguments:
CC  SYSFIL  =      name of system file containing the updated
CC                      file names. The file name in this system
CC                      file will be used by the calling program
CC                      in its subsequent OPEN statement.
CC                      Example: 'USERSYS:ACAxXX.SYS'  where
CC                      USERSYS = logic name of system file directory
CC                      AC      = identifier for calling program ACquire
CC                      A      = machine identifier VAX I (B for VAX II)
CC  NBYT    =      length of system file name in bytes
CC  LSYS    =      log. unit to temporarily open system file
CC  INCR    =      increment by which numeric field of file name is
CC                      modified (may be positive, zero or negative)
CC
CC  return arguments:
CC  FILNAM  =      updated filename to be used in OPEN statement
CC
CC*****
```

BKLIB

BKUTY

LINEPLOT

```
CC*****
CC
CC  F I L E :   LINEPLOT.FOR          PLOT ON LINE-PRINTER
CC  =====
CC                      W. HUERLIMANN
CC
CC*****
CC
```

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC SUBROUTINE LINPLT(XEX,YEX,SIGEX,NEX,XTH,YTH,NTH,NSIZE,SYMBOL,
CC      *
CC      THEORY,TEXTX,TEXTY1,TEXTY2)
CC
CC NTH>0 : PLOT WITH GIVEN THEORETICAL POINTS
CC NTH=0 : NO THEORETICAL POINTS
CC NTH<0 : CALCULATE THEORETICAL POINTS USING REAL*4 FUNCTION THEORY
CC      (TO BE DECLARED EXTERNAL IN CALLING PROGRAM)
CC
CC COMMON /INOUT/ NIN,NOUT
CC
CC *****
```

BKLIB

BKUTY

BK\_R\_SIN

```
CC
CC FILE: BKRSIN.FOR      Read singles spectra (arrays) in
CC =====             Big Karl data format (ACQUIRE)
CC                       J.Meissburger 2-FEB-83
CC                       B.BRINKMOELLER 1-MAR-1985
CC 15-DEC-1985 B.Brinkmoeller :
CC changed to support REAL*4 data type and 2-dim arrays
CC
CC
C  SUBROUTINE BK_R_SIN (LUN, FILENAME, ID,
C      *
C      MAX_NO_ARRAYS, MAX_LEN_ARRAYS, MAX_NO_COND,
C      *
C      NO_OF_ARRAYS, ARRAY_NUMBER, ARRAY,
C      *
C      TYPE, PARAM, SIZE, GAIN, OFFSET,
C      *
C      PARAM2, SIZE2, GAIN2, OFFSET2,
C      *
C      CONDITIONS, C_PAR, C_LOL, C_HIL,
```



# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```

C      *                                DAYTIME, LCOMM, COMMENT, IFLAG)
CC
CC Input arguments:
C  INTEGER*2 LUN                !      Fortran logical unit number
C  CHARACTER*(*) FILENAME      !      name of file to read
C  INTEGER*2 ID                 !      typ of data desired
C  !      if ID = 'RE','LZ','PL' the routine will read all 1-dim
C  !      arrays and convert them to R*4, I*4 or I*2
C  !      if ID = '2R','2L','2D' the routine will read all
C  !      low resolution (R*4,I*4,I*2) 2-dim arrays
C  !      and convert them to R*4, I*4 or I*2
C  !      if ID = '28','24','22','21' the routine will read all
C  !      high resolution (8Bit, 4Bit, 2Bit, 1Bit)
C  !      2-dim arrays without conversion
C  !      if ID = '2 ' the routine will read all 2-dim arrays
C  !      without conversion
C
C  INTEGER*2 MAX_NO_ARRAYS !      maximum number of arrays
C      INTEGER*4 MAX_LEN_ARRAYS!      maximum length of arrays
C      !      measured in longwords
C  INTEGER*2 MAX_NO_COND  !      maximum number of conditions
CC
CC Input and/or return arguments:
C  INTEGER*2 NO_OF_ARRAYS !      total number of arrays found /
C      !      number of arrays to read
C      !      0 => read all arrays
C  INTEGER*2 ARRAY_NUMBER(MAX_NO_ARRAYS)
C      !      array numbers desired / found
C  INTEGER*2 LCOMM        !      maximum length of comment to be
C      !      passed / length of comment passed
CC
CC Return arguments:
C  INTEGER*4 ARRAY(MAX_LEN_ARRAYS,MAX_NO_ARRAYS)
C      !      array channel content
C  INTEGER*2 TYPE  (MAX_NO_ARRAYS) !      array ID as passed

```

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```

C
C  INTEGER*2 PARAM  (MAX_NO_ARRAYS) !      parameter number (ADC)
C  INTEGER*2 SIZE   (MAX_NO_ARRAYS) !      array size (channels)
C  INTEGER*2 GAIN    (MAX_NO_ARRAYS) !      ADC gain
C  INTEGER*2 OFFSET (MAX_NO_ARRAYS) !      channel offset (origin)
C
C  INTEGER*2 PARAM2 (MAX_NO_ARRAYS) !      parameter number (ADC)
C  INTEGER*2 SIZE2  (MAX_NO_ARRAYS) !      array size (channels)
C  INTEGER*2 GAIN2   (MAX_NO_ARRAYS) !      ADC gain
C  INTEGER*2 OFFSET2(MAX_NO_ARRAYS) !      channel offset (origin)
C                                     !      only used for 2-dim arrays
C
C  INTEGER*2 CONDITIONS(MAX_NO_ARRAYS)
C                                     !      number of conditions (gates)
C  INTEGER*2 C_PAR(MAX_NO_COND,MAX_NO_ARRAYS)
C                                     !      condition parameter (up to 16)
C  INTEGER*2 C_LOL(MAX_NO_COND,MAX_NO_ARRAYS)
C                                     !      low limit channel number
C  INTEGER*2 C_HIL(MAX_NO_COND,MAX_NO_ARRAYS)
C                                     !      high limit channel number
C
C  BYTE DAYTIME      (24,MAX_NO_ARRAYS) !      time of data generation
C  BYTE COMMENT      (LCOMM) !      comment found in block with id 'TX'
C  INTEGER*4 IFLAG    !      file read return status
C                                     ! -3  wrong input parameter list
C                                     ! -2  internal file structure error
C                                     ! -1  end of file during read
C                                     !  0  no error
C                                     !  1  at least one array truncated
CC
CC Common Block for the free style I/O routines
C  COMMON /INOUT/ NIN, NOUT, BATCHI, NBATCHI, BATCHO, NBATCHO
C  INTEGER*4 NIN, NOUT, NBATCHI, NBATCHO
C  LOGICAL*1 BATCHI, BATCHO
CC

```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

CC \*\*\*\*\*

BKLIB

BKUTY

BK\_W\_SIN

CC

CC FILE: BKWSIN.FOR Write singles spectra (arrays) in

CC ===== Big Karl data format (ACQUIRE)

CC B.BRINKMOELLER 1-MAR-1985

CC 17-DEC-1985 B.brinkmoeller :

CC changed to support new (I\*2, Real\*4) data types

CC

C SUBROUTINE BK\_W\_SIN (LUN, FILENAME,

C \* MAX\_NO\_ARRAYS, MAX\_LEN\_ARRAYS, MAX\_NO\_COND,

C \* NO\_OF\_ARRAYS, ARRAY\_NUMBER, ARRAY,

C \* TYPE, PARAM, SIZE, GAIN, OFFSET,

C \* PARAM2, SIZE2, GAIN2, OFFSET2,

C \* CONDITIONS, C\_PAR, C\_LOL, C\_HIL,

C \* LCOMM, COMMENT, IFLAG)

CC

CC The output of this routin is a file in standart BIG KARL format with

CC 1) a beginning block. (ID = 'BG')

CC 2) an optional block for the comment. (ID = 'TX')

CC 3) for each array one describtion block containing information passed

CC in PARAM, SIZE, GAIN, OFFSET, PARAM2, SIZE2, GAIN2, OFFSET2, CONDITIONS,

CC C\_PAR, C\_LOL, and C\_HIL, and data blocks for the ARRAY. (ID = TYPE)

CC 4) an end block. (ID= 'BG')

CC

CC !!! The validity of the data typ is not checked. Make sure to pass the

CC correct TYPE for the array. supported are :

CC TYPE = 'RE' for 1-dim REAL\*4 arrays

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```

CC      = 'LZ' for 1-dim INTEGER*4 arrays
CC      = 'PL' for 1-dim INTEGER*2 arrays
CC      = '2R' for 2-dim REAL*4 arrays
CC      = '2L' for 2-dim INTEGER*4 arrays
CC      = '2D' for 2-dim INTEGER*2 arrays
CC      = '28' for 2-dim 8 Bit arrays
CC      = '24' for 2-dim 4 Bit arrays
CC      = '22' for 2-dim 2 Bit arrays
CC      = '21' for 2-dim 1 Bit arrays
CC
CC Input arguments:
C  INTEGER*2 LUN           !      Fortran logical unit number
C  CHARACTER*(*) FILENAME !      name of file to write
C  INTEGER*2 MAX_NO_ARRAYS !      maximum number of arrays
C  INTEGER*4 MAX_LEN_ARRAYS!      maximum length of arrays
C                               !      measured in long words
C  INTEGER*2 MAX_NO_COND  !      maximum number of conditions
C
C  INTEGER*2 NO_OF_ARRAYS !      number of arrays to write
C  ! NO_OF_ARRAYS is not necessarily identical to MAX_NO_ARRAYS.
C  ! So there is a possibility to skip arrays in the input list by setting
C  ! ARRAY_NUMBER of the array to zero.
C
C  INTEGER*2 ARRAY_NUMBER(MAX_NO_ARRAYS)
C                               !      array numbers to write
C  INTEGER*4 ARRAY(MAX_LEN_ARRAYS,MAX_NO_ARRAYS)
C                               !      array channel content
C
C  INTEGER*2 TYPE  (MAX_NO_ARRAYS) ! array identifier
C
C  INTEGER*2 PARAM  (MAX_NO_ARRAYS) !      parameter number (ADC)
C  INTEGER*2 SIZE   (MAX_NO_ARRAYS) !      array size (channels)
C  INTEGER*2 GAIN    (MAX_NO_ARRAYS) !      ADC gain
C  INTEGER*2 OFFSET  (MAX_NO_ARRAYS) !      channel offset (origin)
C

```

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```

C  INTEGER*2 PARAM2 (MAX_NO_ARRAYS) !      parameter number (ADC)
C  INTEGER*2 SIZE2  (MAX_NO_ARRAYS) !      array size (channels)
C  INTEGER*2 GAIN2  (MAX_NO_ARRAYS) !      ADC gain
C  INTEGER*2 OFFSET2(MAX_NO_ARRAYS) !      channel offset (origin)
C                                     !      only used for 2-dim arrays
C
C  INTEGER*2 CONDITIONS(MAX_NO_ARRAYS) !    number of conditions (gates)
C  INTEGER*2 C_PAR(MAX_NO_COND,MAX_NO_ARRAYS)
C                                     !    condition parameter (up to 16)
C  INTEGER*2 C_LOL(MAX_NO_COND,MAX_NO_ARRAYS)
C                                     !    low limit channel number
C  INTEGER*2 C_HIL(MAX_NO_COND,MAX_NO_ARRAYS)
C                                     !    high limit channel number
C
C  INTEGER*2 LCOMM      !      length of comment in byte
C  BYTE COMMENT(LCOMM) !      comment
CC
CC return argument
C  INTEGER*4 IFLAG      !      file write return status
C                                     ! -2  error in input list
C                                     ! -1  error code during write
C                                     !  0  no error
CC
CC Common Block for the free style I/O routines
C  COMMON /INOUT/ NIN, NOUT, BATCHI, NBATCHI, BATCHO, NBATCHO
C  INTEGER*4 NIN, NOUT, NBATCHI, NBATCHO
C  LOGICAL*1 BATCHI, BATCHO
CC
CC *****

```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKLIB

BKUTY

BK\_WAIT

```
C;
C;
C;  FILE: BKWAIT.MAR          Process wait utility
C;  =====                 R.Weber 1-FEB-80
C;                             J.Meissburger 25-JAN-83
C;
C;  FORTRAN SUBROUTINE CALL:
C;
C;  CALL BK_WAIT (absolute_time)
C;  *****
C;
C;  where:  absolute_time = character string specifying
C;                             absolute time
C;                             (see HELP SPECIFY ABSOLUTE)
C;
C;  e.g.:          CALL BK_WAIT ('0 00:00:05.00')
C;
C;          to hibernate process for 5 seconds
C;
C;  *****
C;
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKLIB

BKUTY

BK\_TT\_MODE

CC

CC

CC FILE: BKTTMODE.FOR Set terminal mode (device independent)

CC ===== J.Meissburger 7-Jan-1986

CC

CC VERS: 31-JAN-1986

CC

C SUBROUTINE BK\_TT\_MODE (MODE, LOG\_UNIT, IFLAG)

CC

C CHARACTER\*(\*) MODE

C INTEGER\*4 LOG\_UNIT

C INTEGER\*2 IFLAG

CC

CC Input arguments:

CC

CC MODE = character string symbol specifying the desired  
terminal mode (e.g. 'TCS\_MODE' or 'DEC\_MODE')

CC or

CC graphics enhancement (e.g. 'G\_LINestyle\_5')

CC

CC LOG\_UNIT = logical unit for configuration write

CC

CC Output arguments:

CC

CC IFLAG = 0 for success

CC -1 for configuration failure (any reason)

CC

CC Remark: The mode symbols correspond to the logical file names used

CC in BK\_TT\_CONFIG, but not all (special) features are

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC          generally available. The most important symbols are:
CC          'TCS_MODE' to switch a terminal in PLOT10 compatibility mode
CC          'DEC_MODE' to switch to DEC/ANSI mode (EDIT-mode)
CC
CC *****
```

BKLIB

BKUTY

BK\_DECODE

```
C;
C;
C;
C; FILE: BKDECODE.MAR      DECODE A SINGLE EVENT INTO ADC'S
C; =====              AUTHOR: H.Stoff, ZEL
C;                        J.MEISSBURGER 13-FEB-85 modified
C;
C .TITLE BK_DECODE        Decode a single Memphis event
C;
C;
C; FORTRAN SUBROUTINE CALL:
C;
C; CALL BK_DECODE (LISTMODE_EVENT,
C; 1              BIT_POSITION_TABLE,
C; 2              BYTE_POSITION_TABLE,
C; 3              WORD_LENGTH_TABLE )
C;
C;
C;
C; n      PARAMETER LIST:      n(AP) = addr. of argument
C;                        @n(AP) = argument (immediate)
C;
```



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
C; +
C; 0 NO. OF ARGUMENTS = 4
C;
C;
C; 4 LISTMODE_EVENT INTEGER*2 ADDRESS OF EVENT BUFFER
C;
C; 8 BYTE_POSITION_TABLE INTEGER*4 ADDRESS OF BYTE_POSITION_TABLE
C;
C; 12 BIT_POSITION_TABLE INTEGER*4 ADDRESS OF BIT_POSITION_TABLE
C;
C; 16 WORD_LENGTH_TABLE LOGICAL*1 ADDRESS OF WORD_LENGTH_TABLE
C;
C;
C; *****
C;
```

BKLIB

BKUTY

BK\_TT\_CONFIG

CC

CC

CC FILE: BKTTCONF.FOR CONFIGURATION UTILITY

CC ===== J.Meissburger 26-Sep-1985

CC

CC Vers: 28-Jan-1986

CC

CC Subroutine call:

CC

C SUBROUTINE BK\_TT\_CONFIG (CONFIGS, LOG\_UNIT, IFLAG)

CC

C CHARACTER\*(\*) CONFIGS

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
C  INTEGER*4      LOG_UNIT
C  INTEGER*2      IFLAG
CC
CC  The subroutine BK_TT_CONFIG reads the files defined
CC  by "CONFIGS" and copies them to the device defined
CC  opened on logical unit "LOG_UNIT".
CC
CC  Input arguments:
CC
CC  CONFIGS =      one or more logical names or
CC                  file names to be copied,
CC                  e.g. 'ft5$hp_mode,ft5$aids_off'
CC
CC  LOG_UNIT      =      output logical unit
CC                  (Default = 6)
CC
CC  IFLAG          =      error flag 0 = success
CC                  -1 = failure
CC
CC *****
```

BKLIB

BKUTY

BK\_TT\_AST

```
CC
CC
CC  SUBROUTINE BK_TT_AST (DUMMY)      Terminal Ctrl-Y AST routine
CC  ===== J.Meissburger 26-Sep-1985
CC
CC  Subroutine to handle terminal Ctrl-Y
CC
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

CC Input argument:

CC

CC DUMMY = dummy argument, but must be provided in  
CC any call to BK\_TT\_AST

CC

CC Function value returned:

CC

CC BK\_TT\_AST = 0 (FALSE), if no Ctrl-Y was hit  
CC = 1 (TRUE), if Ctrl\_Y was hit

CC

CC -----

CC

CC SUBROUTINE BK\_TT\_AST\_ENA

CC

CC enables Ctrl-Y recognition by program, disables recognition by DCL

CC

CC SUBROUTINE BK\_TT\_AST\_DIS

CC

CC disables Ctrl-Y recognition by program, reenables recognition by DCL

CC

CC Rem: These routines have to be called only o n c e at program

CC initialization and at program exit resp.

CC

CC ----

CC

CC

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKLIB

BKUTY

BK\_GET\_PAR

CC

CC

CC FILE: BKGETPAR.FOR           Get input parameters from foreign command

CC =====                   J.Meissburger 3-Jun-1985

CC

CC VERS: 22-JAN-1986

CC

C   SUBROUTINE BK\_GET\_PAR (PAR, L\_PAR, N\_PAR)

CC

CC The subroutine reads a foreign command parameter list and  
CC returns the number of parameters found, the parameters and  
CC their true string length.

CC

CC Input arguments:

CC

CC none, parameter list read from SYSS\$INPUT

CC

CC Output arguments:

CC

CC CHARACTER\*64 PAR(8) = parameter values returned

CC       INTEGER\*2 L\_PAR(8) = length of parameter strings

CC       INTEGER\*2 N\_PAR    = number of parameters found

CC

CC \*\*\*\*\*

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKLIB

BKUTY

BK\_TRNLOG

CC

CC FILE: BKTRNLOG.FOR            translate logical name to physical device name

CC =====                      J.Meissburger 29-Nov-1985

CC

CC VERS: 5-FEB-1986

CC

C    SUBROUTINE BK\_TRNLOG ( LOG\_NAME, DEV\_NAME, L\_DEV\_NAME )

CC

C    CHARACTER\*(\*) LOG\_NAME, DEV\_NAME

C    INTEGER\*2 L\_DEV\_NAME

CC

CC Input arguments:

CC

CC CHARACTER\*(\*) LOG\_NAME    =            logical name to translate

CC

CC Output arguments:

CC

CC    CHARACTER\*(\*) DEV\_NAME            =            resulting physical device name

CC    INTEGER\*2        L\_DEV\_NAME        =            number of characters in DEV\_NAME

CC

CC Rem: The subroutine translates until there is no further translation.

CC    Special (invisible) leading characters (<ESC>...) are removed so

CC    that the result can be used for string comparison!

CC

CC \*\*\*\*\*

BIB KARL magnet spectrometer operating system HELP LIBRARY  
Author: J.Meissburger

BKLIB

BKUTY

BK\_TT\_TYPE

```
CC
CC  FILE: BKTTYPE.FOR           Get terminal type symbols
CC  =====                   J.Meissburger 13-Jun-1985
CC
CC Vers: 18-APR-1986 (VERS: VMS 4.3)
CC
C   SUBROUTINE BK_TT_TYPE ( LOG_NAME, TT_TYPE, L_TT_TYPE,
C   *                      TT_SYS_TYPE, L_SYS_TYPE )
CC
C   CHARACTER*(*)  LOG_NAME, TT_TYPE, TT_SYS_TYPE
C   INTEGER*2      L_TT_TYPE, L_SYS_TYPE
CC
CC  This subroutine returns the device class or the terminal type
CC  for terminal-class devices (secondary characteristics) for
CC  the device specified by the logical or physical device name
CC  LOG_NAME. Default for LOG_NAME is SYS$COMMAND. TT_TYPE and
CC  TT_SYS_TYPE are set to 'UNKNOWN' if the device (class) is not
CC  known to the system.
CC
CC  For a process running in BATCH mode TT_TYPE = 'BATCH'
CC  will be returned.
CC
CC *****
CC
CC  Input arguments:
CC
CC  LOG_NAME      =      logical device name (default = 'sys$command')
CC
CC  Output arguments:
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC
CC  TT_TYPE      = device class or terminal type for terminal-class
CC                  e.g. "DISK" or "BATCH" or "FT5"
CC  TT_SYS_TYPE = permanent system terminal type
CC
CC  L TT_TYPE    = length of terminal type return string
CC
CC  L SYS_TYPE   = length of permanent type return string
CC
CC *****
```

BKLIB

BKMAT

```
*****
*  Library: B K M A T  *
*****
```

The following modules are available in BKLIB:BKMAT/LIBR  
(All subroutines are part of MATRIX, Author: W.Huerlimann)

```
CHISQR      perform a chisquare fit
CHIPOL      polynomial fit using CHISQR
```

ALGEBRA matrix operations:

```
MTXADD
MTX EQU
MTXINV
MTXMAT
MTXMBT
MTXMLT
MTXMSC
MTXSUB
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

MTXTRA

MTXTRP

MTXUNT

MTXWRT

BKLIB

BKMAT

ALGEBRA

CC\*\*\*\*\*

CC

CC F I L E : ALGEBRA.FOR MATRIX ALGEBRA SUBROUTINES.

CC ===== W. HUERLIMANN 18-OCT-81

CC

CC\*\*\*\*\*

CC

CC F I L E : MTXTRA.MAT COPY MATRICES

CC ===== W. HUERLIMANN APR-81

CC

CC SUBROUTINE MTXTRA(A,R,M,N)

CC

CC COPY THE (M\*N) MATRIX A INTO R.

CC LIT.: BRANDT "DATENANALYSE".

CC

CC

CC F I L E : MTXMSC.MAT MULTIPLY MATRIX WITH SCALAR

CC ===== W. HUERLIMANN APR-81

CC

CC SUBROUTINE MTXMSC(A,R,S,M,N)

CC

CC MULTIPLY THE (M\*N) MATRIX A WITH SCALAR S AND STORE RESULT IN R.



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC LIT.: BRANDT "DATENANALYSE".
CC
CC
CC F I L E :   MTXUNT.MAT           GENERATE UNIT MATRIX
CC =====
CC                               W. HUERLIMANN  APR-81
CC
CC SUBROUTINE MTXUNT(R,N)
CC
CC GENERATE THE (N*N) UNIT MATRIX R.
CC LIT.: BRANDT "DATENANALYSE".
CC
CC
CC F I L E :   MTXTRP.MAT          TRANSPOSE MATRIX
CC =====
CC                               W. HUERLIMANN  APR-81
CC
CC SUBROUTINE MTXTRP(A,R,M,N)
CC
CC TRANSPOSE THE (M*N) MATRIX A AND STORE RESULT IN R.
CC LIT.: BRANDT "DATENANALYSE".
CC
CC
CC F I L E :   MTXADD.MAT          ADD MATRICES
CC =====
CC                               W. HUERLIMANN  APR-81
CC
CC SUBROUTINE MTXADD(A,B,R,M,N)
CC
CC ADD THE TWO (M*N) MATRICES A,B AND STORE RESULT IN R.
CC LIT.: BRANDT "DATENANALYSE"
CC
CC
CC F I L E :   MTXSUB.MAT          SUBTRACT MATRICES
CC =====
CC                               W. HUERLIMANN  APR-81
CC
CC SUBROUTINE MTXSUB(A,B,R,M,N)
CC
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```

CC  SUBTRACT THE (M*N) MATRIX B FROM THE MATRIX A AND STORE RESULT IN R.
CC  LIT.: BRANDT "DATENANALYSE".
CC
CC
CC  F I L E :   MTXMLT.MAT           MULTIPLY MATRICES
CC  =====
CC                                W. HUERLIMANN  APR-81
CC
CC  SUBROUTINE MTXMLT(A,B,R,M,L,N)
CC
CC  MULTIPLY THE (M*L) MATRIX A WITH THE (L*N) MATRIX B
CC  AND STORE RESULT IN (M*N) MATRIX R.
CC  LIT.: BRANDT "DATENANALYSE".
CC
CC
CC  F I L E :   MTXMAT.MAT           MULTIPLY MATRICES
CC  =====
CC                                W. HUERLIMANN  APR-81
CC
CC  SUBROUTINE MTXMAT(A,B,R,M,L,N)
CC
CC  MULTIPLY THE TRANSPOSE OF THE (L*M) MATRIX A WITH THE (L*N) MATRIX B
CC  AND STORE RESULT IN (M*N) MATRIX R.
CC  LIT.: BRANDT "DATENANALYSE".
CC
CC
CC  F I L E :   MTXMBT.MAT           MULTIPLY MATRICES
CC  =====
CC                                W. HUERLIMANN  APR-81
CC
CC  SUBROUTINE MTXMBT(A,B,R,M,L,N)
CC
CC  MULTIPLY THE (M*L) MATRIX A WITH THE TRANSPOSE OF THE (N*L) MATRIX B
CC  AND STORE RESULT IN (M*N) MATRIX R.
CC  LIT.: BRANDT "DATENANALYSE".
CC
CC
CC  F I L E :   MTXEQU.MAT           SOLVE LINEAR EQUATION

```

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```

CC ===== W. HUERLIMANN APR-81
CC
CC SUBROUTINE MTXEQU(A,B,N,M,IERR)
CC
CC SOLVE THE LINEAR EQUATION A*X=B USING GAUSSIAN ALGORITHM.
CC DIMENSIONS OF MATRICES: A(N*N), X(N*M), B(N*M).
CC THE MATRIX A WILL BE DESTROYED AND THE SOLUTION FOR X
CC IS WRITTEN INTO MATRIX B.
CC LIT.: BRANDT "DATENANALYSE"
CC
CC ERROR FLAG: IERR=0 ALL OK
CC IERR=-1 ERROR
CC
CC
CC F I L E : MTXINV.MAT MATRIX INVERSION
CC ===== W. HUERLIMANN APR-81
CC
CC SUBROUTINE MTXINV(A,R,N,IERR)
CC
CC INVERT THE (N*N) MATRIX A AND STORE RESULT IN R.
CC THE INPUT MATRIX A WILL BE DESTROYED.
CC LIT.: BRANDT "DATENANALYSE".
CC
CC ERROR FLAG: IERR=0 ALL OK
CC IERR=-1 ERROR IN SUBROUTINE MTXEQU
CC
CC
CC F I L E : MTXWRT.MAT PRINT MATRIX
CC ===== W. HUERLIMANN JUN-82
CC
CC SUBROUTINE MTXWRT(A,M,N)
CC
CC PRINT THE (M*N) MATRIX A ON LOGICAL UNIT NOUT.
CC LIT.: BRANDT "DATENANALYSE".
CC

```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

CC \*\*\*\*\*

BKLIB

BKMAT

CHISQUARE

CC\*\*\*\*\*

CC

CC F I L E : CHISQUARE.FOR LEAST SQUARE FIT SUBROUTINES.

CC ===== W. HUERLIMANN 18-OCT-81

CC

CC\*\*\*\*\*

CC

CC

CC F I L E : CHIPOL.FIT POLYNOMIAL FIT USING SUBROUTINE CHISQR.

CC ===== W. HUERLIMANN 25-JUL-81

CC

CC SUBROUTINE CHIPOL(X,Y,SIGMA,NPNTS,COEFF,ERR,NDEG1,IPR,IERR)

CC

CC X = ARRAY (LENGTH=NPNTS) CONTAINING X-VALUES

CC Y = ARRAY (LENGTH=NPNTS) CONTAINING Y-VALUES

CC SIGMA = ARRAY (LENGTH=NPNTS) CONTAINING ERRORS

CC COEFF = ARRAY (LENGTH=NDEG1) OF POLYNOMIAL COEFFICIENTS

CC ERR = ARRAY (LENGTH=NDEG1) OF COEFFICIENT ERRORS

CC NDEG1 = DEGREE OF POLYNOMIAL + 1

CC IPR = PRINTOUT CONTROL : <0 ONLY ERROR MESSAGES ON LUN NOUT

CC =0 SHORT OUTPUT ON LUN NOUT

CC >0 LONG OUTPUT ON LUN NOUT

CC

CC ERROR FLAG: IERR=0 ALL OK

CC IERR=1 ERROR

CC

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC
CC  F I L E :   CHISQR.FIT           GENERAL LEAST SQAURE FIT
CC  =====
CC                               W. HUERLIMANN  25-JUL-81
CC
CC  LIT.: BRANDT "DATENANALYSE"
CC
CC  ERROR FLAG:  IERR=0 ALL OK
CC               IERR=1 ERROR
CC
CC  SUBROUTINE CHISQR(T,Y,CY,N,M,NR,X,CX,ETA,CETA,IPR,IERR)
CC
CC  IMPLICIT REAL*8 (A-H,O-Z)
CC  DIMENSION T(1),Y(1),CY(1),X(1),CX(1),ETA(1),CETA(1),
CC    *          A(2500),B(2500),C(50),XI(50),DELTA(50),EPSLN(50),
CC    *          GB(2500),S1(2500),S2(2500),S3(2500),S4(2500),S5(2500)
CC  COMMON /INOUT/ NIN,NOUT
CC
CC  THE MEASURED VALUES ARE STORED IN Y, THEIR COVARIANCE MATRIX IN CY
CC  AND THE NUMBER OF MEASUREMENTS IN N.
CC
CC  IPR<0 :  PRINT ONLY ERROR MESSAGESON LUN NOUT.
CC  IPR=0 :  SHORT PRINTOUT ON LUN NOUT.
CC  IPR>0 :  LONG PRINTOUT ON LUN NOUT.
CC
CC  *****
CC
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKLIB

AGII

Tektronix Advanced Graphing II

For module description see Tektronix part 4010A02

PLOT10 Advanced Graphing II user manual (Big Karl room 47)

The library BKLIB:AGII.OLB is created with FORTRAN option /I4/NOF77

BKLIB

GMA10

Extended PLOT1 library, but created with FORTRAN option /NOI4 to match  
16 bit DMA access. This library is to be used for access to either  
the GMA102A direct memory access display (Software extension by  
H.Stoff, ZEL) or to a standard TCS compatible graphics terminal with  
automatic terminal configuration and special enhancements  
(Software extension by J.Meissburger using BK\_TT... routines)

For more details see: \$ BKHelp Graphics...

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKLIB

GMAGII

same as AGII, but created with FORTRAN option /NOI4 to match  
the related GMA10 library.

BKLIB

PLOT10

For module description see Tektronix document No. 062-1474-00  
PLOT-10 Terminal Control System users's manual (Big Karl room 47)  
The library BKLIB:PL010.OLB is created with FORTRAN option /I4

BKHELP BRHO (14 July 1986):

BRHO

\*\*\*\*\*

\*\*\* BRHO \*\*\*

\*\*\*\*\*

Author: B.Brinkmoeller

BRHO is Big Karls kinematic program providing the following options:

PLOT ( on graphics line printer or calcomp ploter )

- THE BRHO - VALUE of a particle after a reaction
- THE ENERGY of a particle after a reaction
- THE ENERGY LOSS of a particle in the target  
as a function of
- THE ENERGY OF THE INCOMING PARTICLE
- THE ANGLE OF MEASUREMENT (LAB. ANGLE)

CALCULATE calculate all relevant kinematic data for a reaction as defined  
by an INPUT or SAVE/READ dataset.

The results are stored in a file which may be  
Printed, Typed on your terminal or  
used as an input data file for BKARLO

BRHO

INPUT

With the INPUT command you start entering reaction data.  
The program prompts for the data needed to create a dataset.  
For further use the data may be written to a file BRAXxxx.INP



BIB KARL magnet spectrometer operating system HELP LIBRARY  
Author: J.Meissburger

with the SAVE command.

BRHO

READ

With the READ command you can read data from a previously saved file.

Files with a structure compatible for read are BRAXxxx.INP.  
The program prompts you for the file name.

BRHO

SAVE

With the SAVE command you can write the stored dataset to a file BRAXxx.INP

This dataset can than be restored anytime with the READ command

BRHO

STATUS

Short Information on stored data  
For more Information use the LIST command

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BRHO

LIST

With the LIST command you can examine the stored data. .  
If no further specification is given a summary is displayed.  
To take a closer look some special options are provided.

BRHO

LIST

AL

List all data stored.

BRHO

LIST

RE

List the data of the rest nuclei of the reaction and the  
excitation energies

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BRHO

LIST

TA

List the target data

BRHO

LIST

OU

List the data of the outgoing particle

BRHO

PLIST

Print a list of all data stored on the line printer.

(This command does n o t write the data to a file !)

BRHO

PLOT

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

With the PLOT command you specify the kind of plot you want the program to perform.

The program prompts you for the possible parameters.  
Some values may be omitted to enable defaulting.

If you want to choose the CALCOMP plotter as the output device you have to use the SUBMIT command after you specified the plot. If you don't, the plot will be routed to the graphics line printer after you EXIT the program

BRHO

PLOT

X-SIZE

Length of the frame surrounding the plot in X - direction.  
The default value and maximum is 28 cm.

BRHO

PLOT

Y-SIZE

Length of the frame surrounding the plot in Y - direction.  
The default value is 40 cm.

The maximum value is  $120 * X-SIZE / 28$  cm

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

( for most applications you may simply assume that the  
maximum value is larger than what you need )

BRHO

PLOT

Y-VARIABLE

possible Y-VARIABLES are:

BR	for the BRHO VALUE of a particle after a reaction
E3	for the ENERGY of a particle after a reaction
EL	for the ENERGY LOSS in the target

BRHO

PLOT

Y-LIMITS

You may give minimum and maximum Y - Values to be plotted.  
If you do not enter both, the program will set default values  
so that all output variables belonging to main targets are  
within the frame of the plot.

BRHO

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

PLOT

X-VARIABLE

possible X- VARIABLES are:

TH FOR THE ANGLE OF MEASUREMENT

EN FOR THE BEAM ENERGY

You have to give minimum and maximum X - Values.

BRHO

PLOT

PARAMETER

When you plot as a function of the beam energy

you have to give a fixed value for the lab - angle

When you plot as a function of lab - angle

you have to give a fixed value for the beam energy

BRHO

SUBMIT

With the SUBMIT command you start a separate process

BRAxxxx in the batch - queue that executes all tasks

you have specified before with the PLOT command.

The program prompts you for an output device.

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

You may choose between line printer and CALCOMP plotter.  
The line printer is the default output device.

If you choose the CALCOMP plotter make sure that it is set  
correctly b e f o r e you submit the job. The job will hang up otherwise.

You may submit as many jobs as you want in one run of the  
program and each job can execute any number of plots.

BRHO

EXIT

Return to DCL - level.

If some plots are not submitted by the SUBMIT command, they will  
be executed n o w by a job BRAXxxx that is submitted to the  
batch - queue. The output device is the line printer.

BRHO

EXAMPLE

---

Typical procedure using Program BRHO

```
BR> IN          ! Start input of reaction data
.              ! The program prompts for the data
.
.
BR> SAVE        ! Optionally save the data to a file BRAXxxx.INP
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
BR> PLOT      ! Enter the characteristics of the plot you want
.            ! to create. The program prompts you for the options
.            ! available
.
BR> SUB CAL    ! The data you entered are passed to a batch job
              ! which does the calculation. The output device
              ! will be the CALCOMP plotter.
BR> CAL        ! Calculate all reaction kinematics for the stored
.            ! dataset and write the results to a file BRAXxxx.DAT
.            ! You may PRINT or TYPE this file to have some numbers
.            ! to take home. (and to be happy!)
BR> EXIT      ! Exit the program
```

For more information about the individual commands use the HELP command

BRHO

#### CALCULATE

With the CALCULATE command you write the results of kinematic calculations for the stored data to a file BRAXxxx.DAT. All calculations are performed by a job BRAXxxx which is automatically submitted by the program. You may choose whether you want the output file to be printed at once

BRHO

ABORT



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

If you abort the jobs that were submitted to the batch - queue there will be files BRAXxxx.COM and BRAXxxx.TMP left on your directory. They are used for communication between BRHO and the programs that are doing the calculation. On normal execution these files are deleted automaticaly if they are no longer needed.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP CERN (14 July 1986):

CERN

To access the CERN library or utility programs just include the command

\$ CERN

in your LOGIN file.

CERN

CERNLIB

The CERNLIB object library is included by specifying

\$ LINK ....., CERNLIB/LIBR, .....

in the program link step. It is the concatenated CERN library and resides on CERN\$COM directory.

CERN

---

PATCHY

The PATCHY utility is automatically made available when issuing the \$ CERN command. For more details see: \$ BKHelp PATCHY

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP CTR (14 July 1986):

CTR

\*\*\*\*\*

\* C O S Y T R I E V E \*

\*\*\*\*\*

Author: J.Meissburger

CTR (= poor man's DTR) serves as information storage and retrieval system. The keywords are user-definable with defaults for use as a literature index system.

CTR uses simple, single character commands and prompts for parameters where required. In order to work properly you must at least have read access to the database and its directory. Temporary files are generated on your private directory and deleted after closing the session.

The basic operation with CTR is to establish a collection of records using a joint keyword search (option "C"ollect).

Example:

Collect Accelerator,resonance and

collects all occurrences of string "Accelerator" AND "resonance" in the same record (database entry). The temporary collection may then be typed ("T") or printed ("P") or temporarily saved ("S") for further processing.

>>> Warning: N e v e r try to even read a database file \*.DBS using an Editor such as EDT - it will permanently damage your database ! (Special commands are available to encode/decode to CTR/EDT format)

## BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

### CTR

#### Database

The file name of a database with extension .DBS preceded by device and directory information where appropriate. (The database has a special and fixed format generated by CTR). If you specify the name of a database that does not exist, the system will create a new database with this name. New records are added by using the Insert option ("I").

Example:                    Database: CERN\$INF:CERNLIB

If you frequently access a given database you may define a corresponding symbol in your private LOGIN file such as:

```
$ LIT*erature := 'CTR usersys:ctrconfig.dfl database_name
```

The database is always sorted according to the first ten characters of the first record field. If the first field contains first and family names the sorting automatically acts on the family name. (If you want special sorting criteria you may do so by including a dummy family name at the beginning). First names are characterized by not more than 2 characters followed by a full stop. Family names are 3 or more characters long.

---

### CTR

#### Configuration

The actual keyword fields are defined in a configuration

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

data file. Default is USERSYS:CTRCONFIG.DFL which configures CTR for literature index storage and retrieval. The name of a user-defined configuration file may be given as first parameter when invoking CTR.

Example:

```
MYCTR:=='CTR my_config.dat my_database.dbs
```

CTR

HELP

Type or print this HELP information (see. BKHelp BKHELP)

CTR

COLLECT

Format:

```
Collect keyword[,...] [matching operator]
```

This command searches through the database for the occurrence of the given keyword(s). If several keywords are specified it searches for a logical match of the keywords using the matching operator. Possible operators are AND (Default), OR, NAND and NOR.

If at least one match could be found it temporarily stores the selected records as a COLLECTION which eventually may be typed or printed.

CTR

TYPE

Type [save\_collection[,...]]

Type the actual temporary or saved collections or the entire database if no collection is present. Typing may be interrupted by <CTRL-Y>.

CTR

PRINT

Print [save\_collection[,...]]

Print the actual temporary or saved collections or the entire database if no collection is present.  
Printing is routed to the desired print device. To abort printing  
type: \$ STOP/ABO print\_device

Print devices are the usual print queues with automatic printer configuration for high quality printouts.

The command may also be used to save permanent backups of collections to a non-printing device (e.g. if you specify MTA1: for print\_device)

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

CTR

## INSERT

Insert new records into the database and sort according to first keyword (family name). First names of max. two characters are separated by full stops. Family names must be at least three characters long (pad with blanks if necessary).

All other fields are optional but should be provided whenever possible. (If you omit one of the fields it is recommended to insert a unique character combination which is easily recognized for later replacement !)

Input of single records may be cancelled by typing <Ctrl-Y>, input is terminated by <Ctrl-Z>.

Example:           Option (...): Insert

```
AUTHORS  : X.Y.Ch.Mueller, X.Meier,
AUTHORS  : A. Li
AUTHORS  : <RET>
TITLE    : Any title
TITLE    : <RET>
REFERENCE: JUEL-Spez xxxx
REFERENCE: <RET>
YEAR     : ??Y??
YEAR     : <RET>
KEYWORDS : Accelerator, cooling
KEYWORDS : <RET>
```

At the end of an Insert session the old database may be replaced by the updated one containing the new records. A purge/keep=2 operation will be performed saving the new and the previous versions. (Purging requires appropriate access rights to the older database versions)

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

CTR

STATUS (?)

Displays status information on temporary and saved  
collections

CTR

MERGE

Format:

MERGE collection[,...]] keyword[,...]] [match]

The Merge command allows to create a new temporary  
collection out of one ore more previously saved collections.  
(Use "?" to display your collection status). The collections are  
referenced by their collection numbers, default match is AND.

Example:

Merge 1,3,4 accelerator,resonance and

CTR

DELETE



## BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

This command allows to delete a collection of records from the database. It first collects the desired records, displays them on the screen and asks for the creation of a new database with the selected records removed. Finally it allows to replace the actual database by the new one and performs a purge/keep=2 saving the new and the previous versions. (Purging requires appropriate access rights to the older database versions)

CTR

\*DECODE

Decode CTR format (\*.DBS) into EDT format (\*.DAT)

CTR

\*ENCODE

Encode EDT format (\*.DAT) into CTR format (\*.DBS)

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP CYCLE (14 July 1986):

CYCLE

\*\*\*\*\*

\* C Y C L E \*

\*\*\*\*\*

Author: J.Meissburger

BIG KARLs spectrometer control utility.

It runs on Big Karls PDP-11/40 RSX-11M system and is used to control and monitor the spectrometer status. Related MCR commands are:

MCR> STATUS      to display actual status outside CYCLE  
MCR> CYSTOP      to stop ongoing CYCLE activities

CYCLE

COMMANDS

F I L E :    CYCLE.INF                    24-OCT-80

=====

#XXX      =      numerical argument (format free)  
"XXX      =      alphanumerical argument (string)  
[..]      =      optional (default)

The commands to CYCLE are automatically written to a log-file CYxxxx.BAT which may in turn be used as an input batch files to CYCLE. The files are regularly saved on tape by the @SAVBATCH procedure.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

CYCLE

HELP

HELP           List all valid commands

CYCLE

EXIT

EXIT           Program exit to monitor (MCR)

CYCLE

COMMENT

COMMENT           ["STRING]

Enter the following text as a comment line

---

CYCLE

IMMEDIATE

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

IMMEDIATE "PSNAME #CURRENT

Immediate power supply setting ( no cycling,  
therefore not recommended for D1 and D2 !)

CYCLE

CYCLE

CYCLE "PSNAME #CURRENT [#SLOPE]

Present parameters for main power supply cycling (trough  
maximum) or set HT and QB powersupplies immediately.  
Current values are given in percent, cycling is  
D I S A B L E D with slope set to zero.  
(DEFAULTS: SLOPE = 0.3%/sec for D1,D2 and  
0.6%/sec for Q1-Q3)

CYCLE

MOVE

MOVE "PSNAME #CURRENT [#SLOPE]

Same as " CYCLE ", but without saturation loop (may be  
used to do minor loops).

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

CYCLE

SCALE

SCALE "DVNAME" #PERCENT [#SLOPE]

Scale power supply currents up or down by a positive or negative value given in percent of the actual value. (All QB and HT power supplies are set immediate, the main power supplies have to be moved using the "GO" command).

Scale pream readings up or down by a value given in percent of the actual reading.

("DVNAME=ALL SCALES D1,D2,PRE1-3 SIMULTANEOUSLY")

CYCLE

GO

GO                    Start moving or cycling

CYCLE

MCYCLE

MCYCLE                    Activate QB manual cycling

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

CYCLE

STATUS

STATUS ["DVNAME"]

Print actual device status

CYCLE

BATCH

BATCH ["FILE"]

Enter BATCH mode; the following commands will  
be read from a BATCH file instead of the terminal.

CYCLE

FINISH

---

FINISH

Leave BATCH mode and route control back to the keyboard

CYCLE

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

PREMA

PREMA [#N] [#T]

List NMR and PREMA values N times in  
T minutes time intervals

CYCLE

SET

SET ["DVNAME] [#PAR1] [#PAR2] ...

Set a device according to the given parameters.  
Where available defaults are taken for omitted  
parameters. (Parameters are positional !)

SLX-SLY	position (MRAD), opening angle (MRAD)
HTD1-HTD2	I0, I1, I2, I3 (% per unit distance )
PRE1-PRE3	Prema reading (MV) by setting the corresponding quadrupole Q1-Q3
FU1-FU4	Fumble probe position
MUX	Multiplexer channels for status

CYCLE

RELEASE

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

RELEASE                      Release all digital control units for manual operation

CYCLE

WAIT

WAIT    [#TIME]              Wait a given time (minutes) before executing  
                              the next command string  
                              (DEFAULT: TIME = 0)

CYCLE

SYMBOLS

>LEGAL PSNAMES:              Q, D, S, K, QB, HTD  
                              Q1-Q3, D1-D2, S1, K1-K24, QB51, QB52, QB61, QB62  
                              HTD1-HTD2

>LEGAL DEVICES:              SL, PRE, NMR, FU, MUX  
                              SL1-SL4, SLX, SLY, PRE1-PRE4, NMR1, FU1-FU4, MUX 1-63  
                              ALL = D1-D2, PRE1-PRE3



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP FUMMEL (14 July 1986):

FUMMEL

\*\*\*\*\*

\* F U M M E L \*

\*\*\*\*\*

Author: J.Meissburger

Big Karl magnetic field measurement utility. Used to measure absolute fields with NMR and field gradients with either NMR or Hall probes.

This program runs on Big Karl's PDP11/40 control computer !

FUMMEL

COMMANDS

Command format conforms to Big Karl standards for free style input.

For command descriptions replace:

#XXX = NUMERICAL ARGUMENT (FREE FORMAT)

"XXX" = ALPHANUMERICAL ARGUMENT (STRING)

[..] = OPTIONAL (DEFAULT)

The commands to FUMMEL are automatically written to a log file (\*.BAT) which together with the data files (\*.DAT) are regularly saved on tape by the procedures @SAVBATCH and @SAVDATA.

FUMMEL

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

HELP

HELP            LIST ALL VALID COMMANDS

FUMMEL

EXIT

EXIT            MOVE FUMBLE PROBES OUT OF THE MAGNET GAP AND  
RELEASE FOR MANUAL CONTROL

FUMMEL

COMMENT

COMMENT    [ "TEXT ]

READ THE TEXT ON THIS LINE AS COMMENT STRING

---

FUMMEL

STATUS

STATUS            DISPLAY THE STATUS OF AN ACTIVATED FUMBLE PROBE

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

FUMMEL

SET

SET "DEV [#XSTART] [#XEND]

SET NEW START/END POSITION AND PREPARE FOR  
NEXT MEASUREMENT (ACTIVATE FUMBLE PROBE) OR  
RESET DEFAULT VALUES WITH "DEV = DEFAULT

FUMMEL

PREMA

PREMA [#N] ["TIME]

PRINT NMR AND PREMA READINGS N TIMES  
IN T MINUTES TIME INTERVALS

FUMMEL

GO

GO START NEXT MEASUREMENT

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

FUMMEL

RELEASE

RELEASE

RELEASE DIGITAL CONTROL FOR MANUAL MODE

FUMMEL

BATCH

BATCH "FILE SWITCH FROM TERMINAL INPUT TO BATCH MODE

FUMMEL

FINISH

FINISH

END BATCH MODE AND SWITCH BACK TO TERMINAL

FUMMEL

---

HARDWARE\_DEVICES

Important: The FUMMEL utility is an interrupt-driven software working with the four PREMA digital voltmeters PRE1 .. PRE4. Even if the digital voltmeter readings are not used the software

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

still needs the interrupts in order to perform properly.

>LEGAL DEVICES:       FU1, FU2 - MAGNET D1 (DOWNSTREAM)  
                  FU3, FU4 - MAGNET D2 (DOWNSTREAM)

>NOTE:            IF ANYTHING SEEMS TO GO WRONG, WAIT AT LEAST  
                  70 SECONDS BEFORE YOU GET NERVEOUS !  
                  (FUMMEL will provide an appropriate timeout  
                  if any of the devices does not respond as  
                  expected !)

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP FUPLOT (14 July 1986):

FUPLOT

\*\*\*\*\*

\* F U P L O T \*

\*\*\*\*\*

Author: J.Meissburger

The utility FUPLOT on Big Karl's PDP11/40 control computer is used to graphically present the measurements made with FUMMEL. Command input conforms to Big Karl standards:

F I L E :    FUPLOT.INF                    29-JUL-81

=====

#XXX	=	NUMERICAL ARGUMENT (FREE FORMAT)
"XXX	=	ALPHANUMERICAL ARGUMENT (STRING)
[..]	=	OPTIONAL (DEFAULT)

FUPLOT

HELP

HELP    LIST ALL VALID COMMANDS

---

FUPLOT

EXIT

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

EXIT PROGRAM EXIT BACK TO THE MONITOR (MCR)

FUPLOT

OPEN

OPEN ["FILENAME, ?]

OPEN A NEW DATA FILE ("FILENAME=DV:[XXX,XXX]FILE.EXT)

OR LIST THE NAME OF AN OPEN FILE (?)

THE LAST MEASUREMENT FILE PRODUCED BY "FUMMEL" IS AUTOMATICALLY OPEN ON PROGRAM ENTRY !

FUPLOT

LAST

LAST OPEN THE LAST DATA FILE PRODUCED BY "FUMMEL"

FUPLOT

CLOSE

CLOSE CLOSE A DATA FILE

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

FUPLOT

SET

SET    [#NVARX]    [#NVARY]

SET THE VARIABLE POSITIONS FOR X AND Y IN THE DATA FILE  
(DEFAULT: #NVARX=2 FOR POSITION, #NVARY=5 FOR NMR)

FUPLOT

REWIND

REWIND            REWIND MAGTAPE

FUPLOT

READ

READ    [#N]

---

READ THE NEXT MEASUREMENT OR MEASUREMENT NUMBER #N  
FROM THE DATA INPUT FILE (DEFAULT: NEXT MEASUREMENT)

FUPLOT



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

PLOT

PLOT PLOT THE LAST MEASUREMENT BROUGHT INTO THE BUFFER  
WITH A READ COMMAND

FUPLOT

FIT

FIT [#XLOW] [#XHIGH] [#XO]

PERFORM A MULTIPOLE ANALYSIS OF THE LAST MEASUREMENT  
IN THE BUFFER. DEFAULT POSITIONS MAY BE OVERWRITTEN  
BY SPECIFYING #XLOW, #XHIGH AND #XO. (DEFAULTS ARE  
SET ACCORDING TO THE FUMBLE PROBE NUMBER)

FUPLOT

SAVE

SAVE SAVE DATA AND FIT RESULT ON FILE FUSAVE.TMP (FORMATTED)

FUPLOT

ALL

BIB KARL magnet spectrometer operating system HELP LIBRARY  
Author: J.Meissburger

ALL IDENTIAL TO "READ FIT PLOT"

FUPLOT

LIST

LIST LIST THE LAST MEASUREMENT IN THE BUFFER.

FUPLOT

CURSOR

CURSOR ["ON, "OFF]

SWITCH THE GRAPHICS CURSOR ON OR OFF

FUPLOT

RES

---

RES HARD RESET OF TERMINAL (HP-TERMINALS ONLY)

FUPLOT

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

VARIABLES

>NOTE:           THE VARIABLES ARE: 1 - TIME IN SECONDS  
                  2 - X-POSITION IN MM  
                  3 - PREMA READING  
                  4 -  $D(PREMA)/DX$   
                  5 - NMR READING

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP GRAPHICS (14 July 1986):

GRAPHICS

The standard graphics libraries are GMAGII and GMA10, the extended derivatives of Tektronix AGII and PLOT10 libraries. The full subset of AGII and PLOT10 calls is contained in these libraries but important features are added:

1) The GMA10 library supports a high speed, high resolution TEK GMA102A DMA display (H.Stoff, ZEL).

2) Features such as automatic (e.g. device-independent) terminal configuration, split-screen capability and display file generation capability are added (J.Meissburger). For more details see:

\$ BKHelp Graphics GMA10...

GRAPHICS

AGII

Tektronix Advanced Graphing II

For module description see Tektronix part 4010A02

PLOT10 Advanced Graphing II user manual (Big Karl room 47)

The library BKLIB:AGII.OLB is created with FORTRAN option /I4/NOF77

GRAPHICS

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

## GMA10

same as PLOT10, but created with FORTRAN option /NOI4 to match 16 bit DMA access. This library is to be used for access to either the GMA102A direct memory access display (Software extension by H.Stoff, ZEL) or to a standard TCS compatible graphics terminal. Software extensions provide control of special GMA102A features and terminal initialisation for Big Karl standard terminals as well as special enhancements for graphics terminals such as the HP2623A orr the TEK41xx family colour graphics terminals. (Software extension by J.Meissburger using BK\_TT... routines)

## GRAPHICS

### GMA10

#### Enhancements

Graphics enhancements such as colour are available on the Tektronix 41xx family terminals. The SAMPLE\_PROGRAM shows the use of GMA10 (PLOT10) calls together with the BK\_TT\_MODE enhancement calls.

CC

CC FILE: TTGMA.FOR                    Test GMA10 library and enhancement calls

CC ===== J.Meissburger, 10-Feb-1986

C

C PROGRAM TTGMA

C

C COMMON /TCSINOUT/ TCSIN, TCSOUT

C INTEGER\*4 TCSIN, TCSOUT

C

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC Initialize GMA10 an switch terminal to TCS mode
C
C  CALL INITT(0)
C
C  CALL ANMODE
C  CALL HOME
C
CC Select graphics text index 5
C
C  CALL BK_TT_MODE('G_GTINDEX_5',TCSOUT,IFLAG)
C  WRITE(TCSOUT,20000) 1
C20000      FORMAT(' (',I1,')ANMODE: this message by graphics'
C      * ' alphamode write ')
C  WRITE(TCSOUT,20001)
C20001      FORMAT('      and continued on second line of graphics area')
C
C
CC Select graphics text index 2
C
C  CALL BK_TT_MODE('G_GTINDEX_2',TCSOUT,IFLAG)
C  CALL MOVABS(100,100)
C  CALL AOUTST(50,'(1)AOUTST: this graphtext message to graphics area')
C
CC Change graphics line index and style
C
C  CALL BK_TT_MODE ('G_LINEINDEX_3',TCSOUT,IFLAG)
C  CALL BK_TT_MODE ('G_LINestyle_3',TCSOUT,IFLAG)
C
CC Draw rectangle
C
C  CALL MOVABS(200,200)
C  CALL DRWREL(100,0)
C  CALL DRWREL(0,100)
C  CALL DRWREL(-100,0)
C  CALL DRWREL(0,-100)
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
C
CC Switch terminal back to dialog ANSI mode
C
C  CALL FINITT(-1,0)
C
C  TYPE *,'(1)FINITT (-1,0) called - this message by TYPE to dialog area'
C
CC Repeat same game with different colours
C
C  CALL RESET
C
C  CALL ANMODE
C  CALL HOME
C
C  CALL BK_TT_MODE ('G_GTINDEX_6',TCSOUT,IFLAG)
C  WRITE(TCSOUT,'(/)')
C  WRITE(TCSOUT,'(/)')
C  WRITE(TCSOUT,20000) 2
C  WRITE(TCSOUT,20001)
C
C  CALL BK_TT_MODE ('G_LINEINDEX_7',TCSOUT,IFLAG)
C  CALL BK_TT_MODE ('G_LINESTYLE_7',TCSOUT,IFLAG)
C  CALL MOVABS(400,400)
C  CALL DRWREL(100,0)
C  CALL DRWREL(0,100)
C  CALL DRWREL(-100,0)
C  CALL DRWREL(0,-100)
C
C  CALL FINITT(0,0)
CC
C  TYPE *,'FINITT (0,0) called - this message should not show up at all C'
C  STOP
C
C  END
```

## GRAPHICS

### GMA10

#### Sample\_procedures

Typical procedures to execute an image using GMA10 would be:

- 1) Procedure to execute a graphics application on either SYS\$COMMAND or an allocated split graphics terminal:

```
$ ! ....
$ ask_again:
$ inquire/nopunct device "Graphics device (<RET> = TT): "
$ if device.eqs."" then device = f$logical("TT")
$ assign/user_mode 'device tcs_output
$ assign/user_mode 'device tcs_input
$ run/nodebug {application program}
```

- 2) Procedure modification to force terminal device output:

```
$ if device.eqs."" then .....
$ if f$locate(device,":").eq.f$length(device) then goto ask_again
```

- 3) Procedure modification to create type-specific/nonspecific display file output:

```
$ .....
$ if device.eqs. ....
$ if f$locate(device,":").eq.f$length(device) then -
    inquire/nopunct tcs_out_type -
    "Terminal type override (<RET> = none): "
$ tcs_in_type = tcs_out_type
```



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

\$ .....

GRAPHICS

GMA10

INITT

call: CALL INITT(0)

The routine INITT initializes graphics I/O on either the GMA102 display or on any standard V24-terminal. All terminals except the GMA perform graphics I/O on logical units

```
COMMON /TCSINOUT/ TCSIN, TCSOUT
INTEGER*2 TCSIN, TCSOUT
```

which are assigned to logical files TCS\_INPUT and TCS\_OUTPUT when calling INITT for the first (and only the first) time.

If the logical files are not explicitly reassigned by the user (e.g. \$ ASSIGN/USER TTA1: TCS\_OUTPUT) they are assigned by the system to "TT" (the users terminal) at login.

If the logical names TCS\_INPUT and TCS\_OUTPUT translate to a known and supported terminal device the configuration of the terminal for graphics I/O ("TCS\_MODE") is transparent and automatic. Supported terminal types (See TT\_TYPE) are:

GMA0	GMA102A DMA display
FT2	HP2648A terminal
FT3	TEK4010/12/14 terminal
VT100	VT100 GB (RETROGRAPHICS)

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

FT4	WESTWARD 1015
FT5	HP2623A with ANSI option
FT6	Tektronix 4105A colour graphics terminal
FT7	Plessey PT-100G high resolution terminal

In order to save graphics screens on data files (display files) including all configuration data the two symbols

TCS\_IN\_TYPE    and  
TCS\_OUT\_TYPE

may be defined before calling INITT to override the true (or unknown) terminal type. See \$ BKHelp AGII Example

## GRAPHICS

GMA10

FINITT

call:    CALL FINITT (ix, iy)

The routine FINITT terminates graphics I/O. It also terminates program execution (Fortran STOP) according to the PLOT10 standard if called with positive argument ix.

For negative ix the cursor is placed to the absolute values of (ix,iy) and the terminal is automatically reset to dialogue-mode (Edit mode or "DEC\_MODE"). The internal graphics status is n o t altered so that additional graphics I/O may continue by calling

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

CALL INITT (0) with a graphics page erase  
or CALL RESET without a graphics page erase

## GRAPHICS

GMA10

## REFRSH

REFRSH ( switch, count, x-cursor, y-cursor )  
- control GMA-Refresh-Mode

switch	INTEGER*2	
= 0	stop refresh mode	
= 1	enable refresh mode to fill refresh buffer	
= 2	send refresh buffer to GMA-driver and start display refresh	
= 3	pick up last cursor coordinates of GMA while refresh is active	
count	INTEGER*2	read only
	current length of used refresh buffer.	
	( return after switch = 2 )	
x-cursor	INTEGER*2	read only
y-cursor	INTEGER*2	read only
	cursor coordinates after last "QIO"	
	( return after switch =3, atherwise unchanged )	

## GRAPHICS

### GMA10

### GMA85

GMA85 ( nchar, buffer )

- send a character string to the microprocessor of the GMA-Interface.  
all TEKTRONIX control codes must be send to the mikroprocessor.  
all ASCII-Characters which are not used in a TEK4014-Terminal will be transmitted to the GMA alphanumeric-terminal.

nchar	INTEGER*2 number of characters to transfer ( max. 100 characters )
buffer	INTEGER*2 buffer address of a FORTRAN array

---

## GRAPHICS

### GMAGII

same as AGII, but created with FORTRAN option /NOI4 to match the related GMA10 library.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

GRAPHICS

PLOT10

For module description see Tektronix document No. 062-1474-00

PLOT-10 Terminal Control System users's manual (Big Karl room 47)

The library BKLIB:PL010.0LB is created with FORTRAN option /I4

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP IGM (14 July 1986):

IGM

\*\*\*\*\*

\* I G M \*

\*\*\*\*\*

Authors: U.Hacker, J.Meissburger

Big Karl Interactive Graphics Monitor

Version: 13-Jun-1986 (BK-VAX)

IGM provides an easy way to produce a graphical representation of one- to three-dimensional data on TEK4010 compatible terminals or CALCOMP compatible hardcopy plotters.

No programming effort is required to display or manipulate graphs on your terminal - IGM has it's own command language which can be run interactively or in batch mode or even called by a program to invoke IGM as subsystem to your application (See BKHELP IGM EXAMPLE). For information on parameters and parameter formats type BKHELP IGM PARAMETERS. In case of problems contact U.Hacker

IGM

Example

---

A typical example for an IGM application would be the plotting of Matrix elements calculated with TRANSPORT: The Transport output files are supposed to be TRFILE.\*

\$ IGM

IGM>sys 'directory trfile\*' ! check for input file

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
IGM>Tra on trfile          ! select Transport input
IGM>Tra b2                 ! select b2 plot
IGM>Lineindex 2 gtindex 5 plot ! select colours
IGM>Line 3 tra b3 plot      ! add b3 in diff. colour
IGM>Hcsize 1 gcopy         ! make small hardcopy
IGM>Exit
```

IGM

LINEINDEX

LINEINDEX index

Switch the colour index for line and frame plotting to the value of "index" (0 .. 7 on TEK4105A terminals)

IGM

GTINDEX

GTINDEX index

---

Switch the colour index for alphanumeric text on your plot to the value of "index" (0 .. 7 on TEK4105A terminals)

IGM

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

## GCOPY

Initiate a graphics screen hardcopy on the TEK4695 colour printer  
(Size of copy dependent on HCSIZE setting)

## IGM

### HCSIZE

HCSIZE size

Select large (size=0) or small (size=1) hardcopy format

## IGM

### SYSTEM

Perform a DCL system command. Examples:

IGM>SYStem 'setansi'

IGM>sys 'edt test.bat'

---

## IGM

### AUTOSCALE



# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

## AUTOSCALE

Automatically scale for the data boundaries

## IGM

### BATCH

BATCH filename

Switch the input from keyboard to batch mode,

i.e. read commands from dataset "filename".

To suppress the terminal echo use the PRINT OFF command.

Back to keyboard input at End Of File or when reading  
an END command from batch file.

To temporarily switch back to terminal input use  
the PAUSE command in your batch file.

i.e BATCH HALLO.DAT

HALLO.DAT contains the commands to be executed

## IGM

### CHANGE

```
      { new_X_value new_Y_value [ new_E_value ] }  
      { X new_value                                     }  
CHANGE pos { Y new_value [ new_E_value ]               }
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

{ D }

CHANGE will change data items in the  
data buffer of IGM at line pos.

D means delete the line.

i.e. CHANGE 11 23.56 65.43

in line 11 in the databuffer X will be replaced by 23.56  
and Y by 65.43

IGM

CLEAR

CLEAR

Clear the terminal screen.

IGM

CONTINUE

CONTINUE

---

Continue after a PAUSE command in batch mode

IGM

DATA

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

DATA { X ! Y ! XY ! XYE }

Enter input data from the terminal.

To stop input enter the END command.

i.e. DATA XY

1.0 1.0

2.0 4.0

3.0 9.0

4.0 16.0

5.0 25.0

END

these 5 points will be plotted

IGM

END

END

Switch control back from batch mode to the keyboard.

Stop data input in DATA mode

IGM

EXIT

EXIT

Exit IGM and return control back to system.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

IGM

HARDCOPY

HARDCOPY

Route last picture to the specified hardcopy device

IGM

HPFUNCTION

HPFUNCTION number

Execute a special function for HP-Terminal

number = 3 "next Page"

IGM

IDENT

IDENT graphic input

Mark details in a graph by using the graphics cursor.

Values for graphic input

M : move beam

D : draw between the last to beam positions

T : write text (defined by TEXT I "text") at cursor position

E : exit ident mode

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

IGM

INIT

INIT

Resets all internal values (dynamic restart)

IGM

INSERT

```
                { new_X_value new_Y_value [ new_E_value ] }  
INSERT      pos      { X new_X_value                               }  
                { Y new_Y_value [ new_E_value ]                     }
```

Insert new data at line pos in the IGM data buffer

i.e. INSERT 354 Y 21.43 3.456

After position 354: Y=21.43 and E=3.456

will be inserted (X=0.0)

---

IGM

INPUT

INPUT filename positiondescription

Input reads the dataset filename and stores the data

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

in the data buffer.

Values for "positiondescription"

X for X-column X-value

Y for Y-column Y-value

E for E-column Error (+- error bar)

S for skip an alphabetic field

D to skip numeric data

i.e INPUT ABC.DAT S X S D S Y

reads ABC.DAT like

....

X-WERT 12.33 Y1-WERT 333.4 Y2-WERT 676.8

X-WERT 33.12 Y1-WERT 22.4 Y2-WERT 44.66

X-WERT 43.78 Y1-WERT 1.22 Y2-WERT 3.4

....

X(1)=12.33 Y(1)=676.8 X(2)=33.12 Y(2)=44.66 and so on

will be in the data buffer

IGM

LIST

LIST [ start-pos [ end-pos ] ]

List data in data buffer from line start-pos to line end-pos

i.e. LIST 30

lists the whole data beginnig with position 30

IGM

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

MAGNIFY

MAGNIFY { graphic input ! [ window xmin xmax ymin ymax ] }

defines the magnification window in the origin graph.

Values for first graphic input

1 : identifies window 1

2 : identifies window 2

3 : identifies window 3

Second graphic input

C : continue

E : erase old definition only

A : activates old definition

direct input needs the window number and the x and y ranges

i.e. MAGNIFY 1 22 33 1 4

will magnify in window 1 the data in range x 22 - 33, y 1 - 4

IGM

PAUSE

PAUSE

Temporarely back from batch mode to terminal input

IGM

PLOT

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

PLOT

Draws the data with the specific settings and texts.

IGM

POINT

POINT [ { SCREEN ! VIRTUEL } ]

To measure the position of points with the cursor in  
screen or virtual coordinates.

The default is VIRTUEL.

IGM

PRINT

PRINT [ { ON ! OFF } ]

Switches the batch mode or input mode echo on or off

The default is ON.

---

IGM

READ

READ filename

Reads data in IGM internal format from filename



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

This data will be produced by the WRITE command.

IGM

RESTORE

RESTORE filename

Restores the plot menu from filename.

The menu will be written by SAVE.

IGM

SAVE

SAVE filename

Writes the plot menu into filename.

Keeps it for later use.

IGM

SCALE

SCALE [FIXED] xmin xmax ymin ymax

Sets the minimum and maximum for the range of  
the axes of a graph.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

i.e. SCALE 1 10 -10 10

Only data in the given range will be drawn.

IGM

SHOW

SHOW

Display the plot menu on the terminal.

IGM

SET

SET parameter

Define a value in the plot menu.

For more information type : HELP SET parameter

IGM

SET

AXIS

SET            AXIS { X ! Y } { TICS ttype [ tnumber ] }

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

{ LABEL lfunction }

ttype Tic type format nm (n minor, m major tics)

value range 1 - 6

default 25

tnumber density of tic marks

value range 1 - 5 without 6 - 10 with minor tics

default 8

lfunction type of axis numbering.

1 linear with description

2 logarithmic

3 linear without axis description

4 logarithmic

For every curve therer is an separate axis

-1 linear normal position

-2 lograithmic

-3 linear opposite position

-4 logarithmic

default 1

IGM

SET

.....  
DEFAULT

SET DEFAULT

Reset all values in the plot menu to the default values.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

IGM

SET

ERROR

SET ERROR errvalue [ ssize ]

Sets errors of Y-Values in percent of actual values.

If errvalue eq. 0. data is taken from E-vector of databuffer.

ssize is the scaling factor for the symbol height.

The symbol type "stype" MUST BE NEGATIVE !!!!

IGM

SET

FRAME

SET FRAME [ { Y ! N } ]

Y a plot frame will be drawn.

N no plot frame will be drawn.

Deault Y.

IGM

SET

LINE

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

## SET LINE ltype

Defines the line type

Range for ltype 1 - 9 and -1, -4, -11

-1 no line

-4 point plot

-11 Histogram with variable bin width.

## IGM

### SET

## MULTILOT

### SET MULTILOT mcount

With mcount = -1 its possible to draw an arbitrary number of graphs in one screen.

## IGM

### SET

## SYMBOL

### SET SYMBOL stype

Defines the symbol at the exact position of the data points

Range for stype is 0 - 11

Default is 0 (No symbol).

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

To mark the Error stype must be negative.

IGM

TEXT

TEXT { X1!X2!Y1!Y2!C1!C2!I!D } 2 text2

Defines the 2 text2 for the text arrays X1 - I

D deletes all definition.

Is 2 text2 missing the old value for the specified field  
is erased.

TEXT POS { graphic input ! [ num xpos ypos ] }

Defines the text position on screen.

Graphic input.

X X-text fields

Y Y-text fields

C C-text fields

E exit graphic input.

Direct input

num is 1 for X, 2 for Y, 3 for C,

xpos,ypos are in screen coordinates.

IGM

TRANSPORT

TRANSPORT { ON ! OFF } filename

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Switches input mode to special for reading TRANSPORT files

TRANSPORT ?

Displays the input file name.

TRANSPORT element

Defines the Matrix element for graph (B1-B6 are the beam)

i.e TRANSPORT R22 and so on.

TRANSPORT ELLIPSE {XT!XP!XD!YT!YP!YD} pos

Defines witch ellipse will be drawn

if pos is negative it will be the position in the data buffer

IGM

WINDOW

WINDOW { graphic input ! [ number xmin xmax ymin ymax ] }

Defines the window on the screen to display the data.

Graphic input values

First input

blank main window

1 magnify window 1

2 magnify window 2

3 magnify window 3

secound input

C continue input

E deletes old coordinates

Direct input Number 0 for main window and so on,

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

xmin-ymax will be in screen coordinates.

IGM

WRITE

WRITE filename

Writes the data buffer into filename for later use  
in internal IGM format.



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP MATRIX (14 July 1986):

MATRIX

\*\*\*\*\*

\* M A T R I X \*

\*\*\*\*\*

Author: W.Huerlimann

The Big Karl spectrometer optimization system.

MATRIX calculates the elements of the BIG KARL TRANSPORT matrix using experimental raytracing. These "aberrations" are determined by least square fitting procedures of the measured data.

The influence of the aberrations to the line broadening may be calculated in order to decide which aberration should be minimized with multipole corrections.

The program reads temporary files generated by ACQUIRE containing peak position data as function of SLX,Y slit positions. This data is analyzed by MATRIX in terms of multipoles depending on slit positions and peak positions along the focal plane (energy)

MATRIX

SUMMARY

=====

How to use M A T R I X

W. Huerlimann 13-Nov-81

=====

# BIB KARL magnet spectrometer operating system HELP LIBRARY.

Author: J.Meissburger

Meaning	Command	System prompts	Commands to change data or parameters
-----			
Start	Username:USER		
	Password:none	\$	
	MATRIX	\$	
	RUN MATRIX	MTX>	
Data set-up	INPUT ipeak	=	
	ACREAD	=	
	CORRECT ipeak	COR>	vname nline value
	CLEAR	MTX>	
	READ filename	MTX>	
	SHIFT	MTX>	
Control outputs	LIST ipeak [ALL]	MTX>	
	PLOT ipeak	MTX>	SET SIZE nsize
	STATUS	MTX>	
Calculate aberrations	FIT ipeak [ALL]	MTX>	SET DEG ndeg
	FIT aberr	MTX>	SET DEG ndeg
Minimize aberrations	MINIMIZE aberr	MIN>	(see HELP MINIMIZE)

## Output of results

- Aberrations:	LIST aberr	MTX>	
	PLOT ipeak	MTX>	SET SIZE nsize
	PLOT aberr	MTX>	SET SIZE nsize
- Broadenings:	BROAD channel [DEF]	MTX>	SET SLX opening
	PLOT BROAD	MTX>	SET SLX opening

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

	PLOT BSL	MTX>	SET SIZE nsize SET SLX opening SET SIZE nsize
Line printer outputs	PBROAD, PLIST	MTX>	as BROAD, LIST
	PLOT, PSTATUS	MTX>	as PLOT, STATUS
--> submit to LP	PRINT	MTX>	

## Specials

- Save data	SAVE filename	MTX>
- Read data	READ filename	MTX>
- Batch mode	BATCH filename	BAT:
- Help	HELP command	MTX>
- Print this file	SHORT	MTX>

Stop	EXIT	\$
------	------	----

```
=====
=====
```

## MATRIX

### PARAMETERS

Parameters	allowed values	Meaning
aberr	R12,T122,...,T1222222	aberrations up to 6th order
channel	any number	channel number
filename	up to 9 characters (without extension)	name of batch or data files defaults:

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

		MATRIX.DAT for READ or SAVE
		MATRIX.BAT for BATCH
ipeak	1,2,...,8	peak number
ndeg	<7 for measurements <4 for aberrations	degree of fit polynomial
nline	1,2,...,15	data line number
nsiz	1,2,3	plot size: small, medium, large
opening	any number >0	half opening of entrance slit
value	any number for BRHO	new value
		POSITION
		THETA
		>0 for SIGMA
		>1 for COUNTS
vname	BRHO	magnetic rigidity (kG*cm)
	THETA	SLX-entrance angle (mrad)
	POSITION	peak position (channel)
	SIGMA	half peak width (channel)
	COUNTS	counts in peak

=====

## MATRIX

### COMMANDS

MATRIX command structure is consistent with Big Karl free style input rules. There are three distinct command levels:

```

MAT>    MATRIX main program level
MIN>    MINIMIZE subprogram level
COR>    CORRECT subprogram level
  
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

MATRIX

COMMANDS

CORRECT

Call the correction subroutine to change the data of one peak.

Format: CORRECT [ipeak vname nline value]

ipeak = peak number to be corrected: 1,2,...,8

vname = variable to be corrected: BRHO

THETA

POSITION

SIGMA

COUNTS

nline = line number (not used if vname=BRHO)

value = new value

Special commands:

CORRECT ipeak DELETE nline      delete one data line

CORRECT ipeak DELETE ALL      delete the whole peak

CORRECT ipeak      set program into correction mode

In the correction mode with prompting COR> only the commands

vname nline value

or DELETE nline [ALL]

are used. Return to MTX> with carriage return <CR>.

Caution: Before using the CORRECT command you should list the data with the LIST command, because the data points always are ordered to descending values of THETA !

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

## MATRIX

### COMMANDS

#### EXIT

Program exit back to VAX monitor \$.

## MATRIX

### COMMANDS

#### FIT

Fit a peak or an angular aberration with a polynomial and calculate the TRANSPORT matrix elements up to the order DEG. The polynomial degree is given by the parameter DEG, which may be changed with the command SET DEG.

Format:        FIT [param]

Parameters:   ipeak = peak number: 1,2,...,8

                 Fit one peak and calculate the angular  
                 aberrations R12, T122 etc.

ALL            Fit all peaks.

aberr = angular aberration: R12,T122,...,T1222222

                 Fit an angular aberration and calculate the  
                 corresponding dispersive aberrations T126,  
                 T1266,...,T1226,T12266,..etc.

## BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

## MATRIX

## COMMANDS

HELP

Explain a specified command.

Format: HELP [command]

MATRIX

## COMMANDS

## INPUT

Data input for a specified peak number. (ipeak=1,2,...,8).

Format: INPUT [ipeak]

In the input mode the program asks for input of the following

```
values:   BRH0(ipeak) = magnetic rigidity (kGauss*cm)
```

(required only for a new peak number)

THETA(n) = SLX-entrance angle (mrad)

POS(n) = peak position (channel)

SIGMA(n) = half peak width (channel)

COUNTS(n) = counts in peak

Each peak number contains one BRH0 value and up to n=15 data

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

points THETA, POS, SIGMA, COUNTS.

Return from input mode to MTX> by typing character E.

MATRIX

COMMANDS

LIST

List data on terminal.

Format: LIST [param]

Parameters: ipeak = peak number: 1,2,...,8

List the data of one peak.

ALL List the data of all peaks.

aberr = angular aberration: R12,T122,...,T1222222.

(if the aberration has been fitted, also the  
corresponding dispersive aberrations will  
be listed.)

MATRIX

COMMANDS

PLOT

Plot on terminal. If the data to be plotted has been fitted, the



## BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

theoretical curve will be displayed with asterisks (\*). The plot size is defined by the parameter SIZE (see command SET SIZE).

Format: PLOT [param]

Parameters: ipeak = peak number: 1,2,...,8

Plot POS vs THETA.

aberr = angular aberration: R12,T122,...,T1222222

Plot aberration vs position.

BRO Plot the theoretical line broadening due to the calculated aberrations as a function of the position in the focal plane. (only fitted aberrations included!)

BSL Plot the theoretical line broadening as a function of of the half slit opening SLX (see command SET SLX).

## MATRIX

### COMMANDS

#### READ

Read the peak data from the specified data file. If no file name is given, the default file MATRIX.DAT will be read in. This command clears the already stored data and calculated aberrations.

Format: READ [filename]

Parameter: filename = file name (up to 9 characters)

If not blank, the file

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

filename.DAT will be read.

## MATRIX

### COMMANDS

#### SAVE

Save all stored peak data to the specifies data file. If no file name is given, the default file MATRIX.DAT is used. This command overwrites the old file.

Format:       SAVE [filename]

Parameter:   filename = file name (up to 9 characters)  
                  If not blank, the data will be  
                  saved into the file filename.DAT

## MATRIX

### COMMANDS

#### SET

Set a specified parameter to a new value.

Format:       SET [param]

Parameters:   CONV = conversion factor between channels and cm.

## BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

DEG = degree of fit polynomial: 0,1,...,6 for meas.,  
0,1,2,3 for aberr.  
RBND = mean bending radius (cm)  
SIZE = size of plots: 1 small plot (60x20 charact),  
2 medium plot (90x40 charact),  
3 large plot (120x60 charact).  
SLX = half slit opening (mrad).  
XMID = channel position for central ray  
(i.e. middle of detector)  
DEF set default values:  
DEG=2, SIZE=1, SLX=10 mrad,  
CONV=48.8 ch/cm, XMID=1000 ch,  
RBND=198.16 cm.

### MATRIX

### COMMANDS

### STATUS

Display the current status of the program. The following informations will be given:

- Values of the parameters DEG,SIZE,SLX,XMID,CONV,RBND  
(to change parameters use command SET)
- Stored data
- Calculated aberrations
- Calculated values of spectrometer dispersion R16, field value B and magnetic rigidity BRHO corresponding to position XMID.

Author: J.Meissburger

## COMMANDS

Calculate and list the theoretical line broadening due to the calculated aberrations at given positions, using the half slit opening SLX ( see command SET SLX ). Only fitted aberrations are included in these calculations.

Parameters:    channel = channel number(s) for which broadening  
                              has to be listed.  
                  DEFAULT    list broadening for the default channel  
                              numbers 0, 200,..., 1800, 2000.

## COMMANDS

Switch program to batch mode and read all commands from the specified batch file. If no file name is given, the default batch file MATRIX.BAT is used.

Parameter:     filename = file name (up to 9 characters)

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

If not blank, the file  
filename.BAT is used.

To return to terminal input, the last command in the batch file  
must be FINISH.

MATRIX

COMMANDS

FINISH

Last command in batch files used to switch back to terminal input  
(see command BATCH).

MATRIX

COMMANDS

SHORT

List a short overview of the MATRIX commands on the line printer.

---

MATRIX

COMMANDS

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

PBROAD

Same as BROAD, but with line printer as output device.

Further explanations: see commands BROAD and PRINT.

MATRIX

COMMANDS

PLIST

Same as LIST, but with line printer as output device.

Further explanations: see commands LIST and PRINT.

MATRIX

COMMANDS

PLOT

Same as PLOT, but with line printer as output device.

Further explanations: see commands PLOT and PRINT.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

MATRIX

COMMANDS

PSTATUS

Same as STATUS, but with line printer as output device.

Further explanations: see commands STATUS and PRINT.

MATRIX

COMMANDS

PRINT

Submit the print output which was generated with the Pcommands (i.e. PBROAD, PLIST, PPLOT, PSTATUS) to the line printer.

This output-file is called MATRIX.OUT and will be erased after printing.

MATRIX

COMMANDS

ACREAD

Read the ACQUIRE file 'ACMTX.TMP'.

## BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Format:       ACREAD [param]

Parameters:   ON     read file from directory [USER.ACQON]  
              OFF    read file from directory [USER.ACQOFF]  
              BACK   read file from directory [USER.ACQBACK]

This file contains the peak integration data items CENT, SIGMA and SUM, which are to be stored into the variables POS, SIGMA and COUNTS. Hereby at most 8 peaks per spectrum are accepted. This new data will be added to already existing data, i.e. 1st data line to peak-nr 1, 2nd data line to peak-nr 2, etc.

If the data of a new spectrum (indicated by a blank line) is encountered, the program prints the centroid value CENT of the the first data line (which helps to identify the spectrum) and asks for the input of the SLX-angle THETA. At most 15 different angles THETA may be defined.

To skip one spectrum type 'S', to terminate type 'E'.

When a peak which was not present before (i.e. empty peak-nr) is encountered, the program prints the value of CENT and asks for the BRHO value of this peak.

To terminate type 'E'.

How to create the file ACMTX.TMP : Use the ACQUIRE command  
                                  'HELP SUM'.

MATRIX

COMMANDS

MINIMIZE



Author: J.Meissburger

MTX> HELP MINIMIZE command

DELETE

Parameter:      imeas = measurement-nr (=line number)

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

MATRIX

COMMANDS

MINIMIZE

EXCLUDE

Exclude one of the stored aberration data points from the FIT and PLOT in the minimization.

Format:            EXCLUDE [imeas]

Parameter:        imeas = measurement-nr (=line number)

MATRIX

COMMANDS

MINIMIZE

INCLUDE

Include an aberration data point in FIT and PLOT for the minimization.

Format:            INCLUDE [imeas]

Parameter:        imeas = measurement-nr (=line number)

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

If a new point is to be included, the program asks for the multipolestrength for this measurement.

MATRIX

COMMANDS

MINIMIZE

RETURN

Return to MTX>

MATRIX

COMMANDS

MINIMIZE

PLOT

Plot the aberration under consideration vs the multipole strength. If the aberration has been fitted (see MINIMIZE-command FIT), the theoretical curve is displayed through asterisks (\*) and the optimal point (if found) is plotted as X.

MATRIX

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

COMMANDS

MINIMIZE

MULTIPOLE

Change the multipole strength for one meas-nr.

Format:           MULTIPOLE [imeas]

Parameter:       imeas = meas-nr (=line number)

MATRIX

COMMANDS

MINIMIZE

LIST

List the aberration data under consideration. The list contains the following items:

MEAS-NR	= measurement number
FILENAME	= last file name given in the SAVE or READ command
DEGREE OF FIT	= degree of fit polynomial from which the aberration has been calculated
MULTIPOLE STRENGTH	= multipole strength (arbitrary units)
ABERRATION	= value of aberration
ERROR	= estimated error of aberration

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

INCLUDED                   = YES if point is included in FIT/PLOT  
                          = NO if point is excluded from FIT/PLOT

MATRIX

COMMANDS

MINIMIZE

FIT

Fit the aberration under consideration as a function of the multipole strength and calculate the optimal multipole setting and the corresponding driving term in the minimum of this curve. The fitting is done with a polynomial of degree DEG (see command SET DEG) which must have the value 1, 2 or 3.

MATRIX

COMMANDS

CLEAR

Clear all stored data and calculated aberrations.  
Clear BRHO-values is optional (CLEAR ALL)

MATRIX

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

## COMMANDS

CLEAR

ALL

Clear BRH0-values as well to measure new spectrum

## MATRIX

### COMMANDS

SHIFT

Shift the stored peak-nrs up by one step, i.e. copy peak-nr 7 into peak-nr 8, copy peak-nr 6 into peak-nr 7, etc. The old data of peak-nr 8 will be destroyed and you gain space in peak-nr 1 in order to read in a new peak with ACREAD.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP MEMPHIS (14 July 1986):

MEMPHIS

\*\*\*\*\*

\*\*\* MEMPHIS \*\*\*

\*\*\*\*\*

Authors: J.Meissburger, R.Korthues

User utility to configure the Big Karl MEMPHIS data acquisition system HARDWARE. Most commonly used to change coincidence conditions or to perform a system soft reset with default configuration loading.

For more details see: HELP EXAMPLE

MEMPHIS

EXAMPLE

- 1) Invoke MEMPHIS at DCL command level by typing:

\$ MEMPHIS

- 2) MEMPHIS first displays the actual hardware status and prompts you for commands with:

MEM>

- 3) You may now change your software status by means of the SET command (See HELP SET). The most important commands are:

MEM> SET COIncidence ## or

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

MEM> SET NOCoincidence ##

which includes an MPS-ADC number ## in the coincidence (default coincidence time is 4.0 usec) or excludes it from coincidence.

I M P O R T A N T :      At present at least one MPS\_ADC has to run in coincidence with itself thus determining implicitly a trigger coincidence with the external event start.

You may check your software status any time with the command:

MEM> SHO SOFt [ALL]

4) If you are satisfied with your software status exit MEMPHIS:

MEM> EXIT

The system then asks you if you really want to download this configuration into the MEMPHIS hardware: If so, respond "Y" and wait a few seconds (especially in case of a RESET) for the system to be configured.

5) The download link terminates operation at

MPSLINK>

command level. Watch a few seconds for ADC or system errors. If everything looks alright, exit the link with:

MPSLINK> EXIT

If you find many ADC errors (error 24) something is probably wrong with either the external event start or the hardware coincidence timing of the ADC's.

6) Your hardware is now ready for data taking. You may check again



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

its actual status with the DCL command:

```
$ MPSTATUS or
$ ACSTATUS
```

You are now ready to run ACON and take data: Good luck !

7) In case of hardware trouble perform a HARD-RESET by typing

```
$ MEMPHIS
MEM> HARD
..... R
MEM> RELOAD
MEM> EXIT
```

The RESTORE command restores the last configuration generated by the previous MEMPHIS session (which usually is your desired experimental configuration)

MEMPHIS

EXIT

Invoke the MEMPHIS microprocessor link or exit back to DCL command level

MEMPHIS

SET

## BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

The SET command presets a device configuration for downloading with MPSLINK. Only the software status of MEMPHIS is modified by this command - hardware status only changes on request after EXItting the MEMPHIS utility !

### MEMPHIS

#### SET

#### COINCIDENCE

SET COINCIDENCE n [time]

Include MPS\_ADC (input unit) number n in the slow hardware coincidence

The optional coincidence time is the time in microseconds after the ADC-RTP pulse MEMPHIS will wait to check if the ADC went busy. (Default = 4.0 microseconds corresponding to the ADC gate time). If the ADC did not go busy after that time MEMPHIS will reject the event and perform a fast ADC reset.

(MPS\_ADC's correspond to the MEMPHIS input units and are counted logically from left to right in the MEMPHIS system crates)

### MEMPHIS

#### SET

#### NOCOINCIDENCE

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

SET NOCoincidence n

Remove MPS\_ADC number n from slow hardware coincidence

>>> At least one MPS\_ADC (input unit) has to be included in <<<

>>> coincidence with itself in MEMPHIS coincidence mode ! <<<

MEMPHIS

SET

MPS\_ADC

SET MPS\_adc n [parameters]

Configure the input/output bit range and position in the  
MEMPHIS event output word (128 bit) for a MPS\_ADC

MEMPHIS

SET

MPS\_ADC

INPUT

SET MPS\_adc n INPut n [parameters]

The INPut parameter defines the input range (in bits) for the  
MPS\_ADC (input unit). This will usually correspond to the output range  
of the connected physical ADC. (Default INPut = 11)

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

MEMPHIS

SET

MPS\_ADC

OUTput

SET MPS\_adc n OUTput n [parameters]

The OUTput parameter defines the output range (in bits) of the MPS\_ADC in the MEMPHIS event output word (128 bits). It is usually smaller than or equal to the input range defined by INPut.

MEMPHIS

SET

MPS\_ADC

POSITION

SET MPS\_adc n POSition n [parameters]

The POSition parameter determines the bit position of the MPS\_ADC in the 128 bits long MEMPHIS event output word. If specified without a numerical value the next free position will be taken.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

MEMPHIS

SET

EVENT\_MODE

SET Event\_mode

Configure MEMPHIS for event ( coincidence ) mode. At least one ADC has to be in coincidence with itself or with the external EVENT START signal in order to define an event. MEMPHIS start may be selected either EXTERNAL by an EVENT START signal or INTERNAL by enabling internal MPS\_ADC start (See HELP SET INTERNAL)

MEMPHIS

SET

SINGLES\_MODE

SET SINGles\_mode [MPS\_ADC #]

Configure MEMPHIS for singles (wired-or) mode. Start mode is automatically set to INTERNAL with all (default) or the desired MPS\_ADC's enabled for starting MEMPHIS.

MEMPHIS

SET

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

INTERNAL\_START

SET Internal\_start [MPS\_ADC #]

Enable MEMPHIS internal start mode. All (default) or the desired MPS\_ADC's may start MEMPHIS to define a possible event. The external EVENT START signal is disabled.

MEMPHIS

SET

EXTERNAL\_START

SET EXternal\_start

Configure MEMPHIS for external EVENT START. All internal MPS\_ADC starts are disabled. MEMPHIS should normally be set to EVENT\_MODE.

MEMPHIS

SET

WINDOW

SET WInDow # parameter

Configure the windows of unit #.

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Parameters are:

ENABLE [#,...] Enable windows. Every two numbers are the limits of one window. Default: the whole window enabled (0-8191).

DISABLE [#,...] Disable windows. Every two numbers are the limits of one window. Default: the whole window disabled (0-8191).

MEMPHIS

SHOW

SHOW parameter

Displays information about the program status

MEMPHIS

SHOW

SOFTWARE

SHOW SOFTWARE [parameter]

Shows the present software status of your MEMPHIS session. This will eventually become the MEMPHIS hardware status if downloaded by MPSLINK after exiting the MEMPHIS software utility.

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Parameters are:

- ALL - Display whole software status.
- STATUS - Display general software status.
- MPS\_ADC [#] - Displays the software status of the MPS\_ADC units.
- WINDOW [#] - Displays the software status of the window units.

MEMPHIS

SHOW

HARDWARE

SHOW HARDware [parameter]

Displays the actual MEMPHIS hardware status.

Parameters are:

- ALL - Display whole hardware status.
- STATUS - Display general hardware status.
- MPS\_ADC [#] - Displays the hardware status of the MPS\_ADC units.
- WINDOW [#] - Displays the hardware status of the window units.

MEMPHIS

RESET



## BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Set up a soft RESET for downloading by MPSLINK. This will reload the default hardware configuration when executed by MPSLINK. A soft reset may be performed without the need to exit ACQUIRE.

### MEMPHIS

#### HARD\_RESET

Set up a hard reset. The MEMPHIS hard reset may be used to unblock the MEMPHIS microprocessor in case of serious troubles. It can only be performed if ACQUIRE is not active ! The hard reset reloads the default configuration as does the soft reset.

### MEMPHIS

#### RELOAD

Reloads the software configuration of the last MEMPHIS session.

### MEMPHIS

#### ENABLE

ENable      [parameter]

Enable use of units, parameters are:

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

WINDOW # [,#] - Enable window unit # for use.

MEMPHIS

DISABLE

DISable [parameter]

Disable use of units, parameters are:

WINDOW # [,#] - Disable window unit # for use.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP MINIMIZE (14 July 1986):

MINIMIZE

MATRIX: MINIMIZE subsystem of MATRIX

MINIMIZE

DELETE

Delete one line from the stored aberration data.

Format: DELETE [imeas]

Parameter: imeas = measurement-nr (=line number)

MINIMIZE

EXCLUDE

Exclude one of the stored aberration data points from the FIT  
and PLOT in the minimization.

Format: EXCLUDE [imeas]

Parameter: imeas = measurement-nr (=line number)

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

MINIMIZE

INCLUDE

Include an aberration data point in FIT and PLOT for the minimization.

Format:           INCLUDE [imeas]

Parameter:       imeas = measurement-nr (=line number)

If a new point is to be included, the program asks for the multipole-strength for this measurement.

MINIMIZE

RETURN

Return to MTX>

MINIMIZE

PLOT

Plot the aberration under consideration vs the multipole strength. If the aberration has been fitted (see MINIMIZE-command FIT), the theoretical curve is displayed through asterisks (\*) and the optimal point (if found) is plotted as X.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

MINIMIZE

MULTIPOLE

Change the multipole strength for one meas-nr.

Format:           MULTIPOLE [imeas]

Parameter:       imeas = meas-nr (=line number)

MINIMIZE

LIST

List the aberration data under consideration. The list contains the following items:

MEAS-NR	= measurement number
FILENAME	= last file name given in the SAVE or READ command
DEGREE OF FIT	= degree of fit polynomial from which the aberration has been calculated
MULTIPOLE STRENGTH	= multipole strength (arbitrary units)
ABERRATION	= value of aberration
ERROR	= estimated error of aberration
INCLUDED	= YES if point is included in FIT/PLOT = NO if point is excluded from FIT/PLOT

## MINIMIZE

### FIT

Fit the aberration under consideration as a function of the multipole strength and calculate the optimal multipole setting and the corresponding driving term in the minimum of this curve.

The fitting is done with a polynomial of degree DEG (see command SET DEG) which must have the value 1, 2 or 3.

BKHELP MINUIT (14 July 1986):

MINUIT

\*\*\*\*\*

\* M I N U I T \*

\*\*\*\*\*

Authors: F.James (CERN-MINUIT)  
J.Meissburger, G.Slakov (MINFIT)

MINUIT is the name of a system of programs developed at CERN. These programs are used to minimize a user defined function; the classic example being the chi-square difference between a set of data points and a theoretical function. The two other important cases which MINUIT can handle are maximum likelihood problems and optimization problems.

The user may start from the beginning and, depending on which of the above tasks he wants to be done, write a program with his own function to be minimized. For the problem of getting a chi square fit between a theoretical function and experimental data, there is a utility called MINFIT which uses MINUIT to minimize the chi-square difference between function and data.

MINUIT is a sophisticated set of programs. To use it, you are advised to consult the MINUIT manual obtainable from CERN computing library.

MINUIT

COMMANDS

MINUIT is told what to do and how to do it by various initializations

## BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

and commands; the initializations define parameters, the commands invoke calls to MINUIT subroutines.

The order of a command sequence is always the same:

The first line is a format-free title, the next lines are parameter definitions, followed by a blank line, which is in turn followed by the commands which tell MINUIT exactly what to do in order to minimize the function.

### MINUIT

#### COMMANDS

#### PARAMETERS

The parameters are defined immediately after the title line.

Each parameter line is formatted as follows:

Columns 1 to 10:	Parameter number as referred to by user.
Columns 11 to 20:	Alphanumeric name for the parameter. (optional)
Columns 21 to 30:	Initial estimate of parameter value.
Columns 31 to 40:	Approximate error of parameter. (if 0 or blank, parameter is constant.)
Columns 41 to 50:	Lower bound on parameter. (if blank, no lower bound)
Columns 51 to 60:	Upper bound on parameter. (if blank, no upper bound)

Input all numbers in real format (ie. 2. as opposed to 2)



# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

## MINUIT

### COMMANDS

#### EXIT

This command indicates that the minimization commands are over, and the user would like to see the results.

## MINUIT

### COMMANDS

#### FIX

This command fixes a parameter at its present value until it is "released" by a RELEASE command. This command might be used if you feel that the given parameter is not allowing the other parameters to minimize properly.

The syntax for this command is:

Column 1 to 3: FIX

Column 11 to 20: Parameter number (use real format, ie 3.0 or just 3.)

## MINUIT

### COMMANDS

## MIGRAD

This command causes the program to perform a minimization using the MIGRAD technique (which is described in the user's manual).

The syntax for this command is:

Column 1 to 6:	MIGRAD
Column 11 to 20:	The maximum number of calls that can be made to the user's function in the MIGRAD minimization. If blank, the default is 1000. (use real format, ie. 500. or 500.0 )
Column 21 to 30:	The tolerance on the value of the minimized function. If blank, the MINUIT assumes a default (See the MINUIT manual for further information.)
Column 31 to 40:	Tolerance on the stability of the error matrix. If blank a default value of 0.01 is assumed. (See the MINUIT manual for further information.)

## MINUIT

### COMMANDS

#### MINIMIZE

This command causes the program to attempt to minimize the users function by first calling SIMPLEX and then MIGRAD.

The syntax for this command is:

Column 1 to 8:	MINIMIZE
----------------	----------

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Author: J.Meissburger

Column 11 to 20: Maximum number of calls to the users function.  
(use real format, ie. 2000.0 or just 2000.)  
If blank, the default is 1000.

Column 21 to 30: tollerance on function value. (default value 0.1)

## MINUIT

### COMMANDS

#### MINOS

This command causes an in-depth error analysis to be done on the present values of the parameters. Because of this, this command is normally only made when an minimum has been found.(ie. after a MINIMIZE, SIMPLEX, or MIGRAD command).

The errors produced are dependent on the uncertainty you have given your input data, so be sure these uncertainties are meaningfull.

The syntax for this command is:

Column 1 to 5: MINOS

Column 11 to 20: The maximum number of calls allowed to your user-defined FCN. (remember to use real format, ie. 800.)

If blank, the default value is 1000.

Column 21 to 80: The parameters that will have MINOS error analysis done to them. The format is every two columns equals one parameter. For example:

12345678901234567890 (These are column numbers, not to be included in file!)

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MINOS      b1b2b612      (where b is a blank)

means that MINOS will be done to parameters  
1,2,6,12 in that order. If columns 21 to 80 are  
left blank, MINOS is done to all parameters.

MINUIT

COMMANDS

RELEASE

This command causes a parameter that has been fixed by a FIX command  
to be allowed to vary again.

The syntax for this command is:

Column 1 to 7:      RELEASE

Column 11 to 20:    The identification number of the parameter to be  
                     released. (use real format, ie. 4.0 or just 4.)

MINUIT

COMMANDS

SIMPLEX

This command causes the program to perform a minimization using a  
technique called the simplex method. See the MINUIT manual for further

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Author: J.Meissburger

information on this technique.

The syntax for this command is:

Column 1 to 7:     SIMPLEX  
Column 11 to 20:   The maximum number of calls permitted to the  
                    user-defined FCN. (use real format, ie 900.)  
                    If blank, the default is 1000.  
Column 21 to 30:   The tolerance on the mininum function value.  
                    If blank, the program assumes a suitable default.

## MINUIT

### COMMANDS

#### EXAMPLE

As an example of what MINUIT can do, let's assume you want to fit a quadratic polynomial to a set of data points.

Your user-defined function would be the sum of the chi-square differences between a set of data points and the quadratic polynomial, defined as

$$\text{theoretical\_function} = \text{PAR}(1) + \text{PAR}(2)*X + \text{PAR}(3)*(X**2).$$

If the following numbers denote column number,  
123456789012345678901234567890123456789012345678901234567890  
then a suitable sequence of instructions might be:

CHI-SQUARE FIT OF A QUADRATIC POLYNOMIAL           (This is the title)

1. OFFSET     45.     1.

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2. LINEAR	.02	.01
3. QUAD	.7	.1

MINIMIZE

EXIT

Notice that all numbers are real-formatted, and that all numbers are in the proper range of columns. It is important to give a reasonable approximation to the expected minimum parameter value and its approximate error, otherwise MINUIT may find a minimum that is physically meaningless, or it may not find a minimum at all.

MINUIT

MINFIT

The MINFIT utility was developed to handle the problem of minimizing the chi square difference between a set of data points and a user defined function .The function actually being minimized is defined as:

$$F = \text{sum of } ((Y_{\text{experimental}} - Y_{\text{theoretical}}) / DY_{\text{experimental}})^2$$

for all values of  $Y_{\text{experimental}}$ , and

where  $DY_{\text{experimental}}$  is the measured variance in the Y values, and  $Y_{\text{theoretical}}$  is the value of the user defined function at a particular X value.

The function to be fitted can be either a polynomial of up to degree 5, or it can be a user defined function of any type.

To begin execution of the MINFIT utility type:

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\$ MINFIT

MINUIT

MINFIT

ORGANIZATION

There are two basic options built into the MINFIT utility; it can either fit a polynomial to your data, or it can fit your user defined function to the data.

If you are fitting your own function to the data, you need two things. One, you need the fortran source for your function, and two, you need to compose a command sequence which will enable MINUIT to fit your function. The Fortran source for your function must be based on the file USERINF:THEORY.FOR, so copy this file into your own directory and modify it into your own THEORY.FOR. The commands to MINUIT are normally placed in a file called COMMAND.DAT. For information on which commands to use, type HELP MINUIT COMMANDS, or see the MINUIT manual.

If you simply want a polynomial fit for your data, then use the polynomial option, described in HELP MINUIT MINFIT POLYNOMIAL.

For information on the various input, output, and fitting options of MINFIT, type BKHELP MINUIT MINFIT INPUT\_FILE.

To execute the MINFIT utility type \$ MINFIT. The resulting interactive procedure will prompt you for all necessary information.

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Author: J.Meissburger

## MINUIT

### MINFIT

#### INPUT\_FILE

The input file contains two things, experimental data and control information for the MINFIT utility.

Default values are supplied for everything except the experimental data, so no command lines are necessary, except of course if you want to change a control or start a new data set. Remember that each data set must be separated by a command line, even if it is only a blank line.

The input file is set up as follows:

#### GLOBAL CONTROL LINES

. . .  
. . .

#### LOCAL CONTROL LINES

. . .  
. . .

#### DATA LINES

. . .  
. . .

#### LOCAL CONTROL LINES

. . .  
. . .

#### DATA LINES

. . .  
. . .

I

I

I===== First Data set

I

I

I

I

I

I===== Second Data set

I

I

I



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ETC.

Any line whose first non-blank character is non-numerical will be interpreted as a command, including completely blank lines.

There are two types of command lines, global and local. A global control will alter that parameter for each data set, whereas a local control will alter that parameter only for the dataset it precedes.

Each control line has the format

CONTROL      [PARAMETER(S)]

Global controls are:

PLOT

By default, no plots are printed. If plots of the theoretical function and of the difference between theoretical points and experimental points are desired, then include this command in the file.

EXTRAPOLATE    [START    FINISH    DELTAX]

Normally the program will only calculate the theoretical function for the experimental values of X. If the user wants the theoretical function evaluated for more or less points than the number of data points EXTRAPOLATE should be used. It calculates the theoretical function at X values between START and FINISH, separated by DELTAX. The command EXTRAPOLATE with no numbers following will cause the function to be evaluated at the following default values:

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START = minimum of Xexp. , FINISH = maximum of Xexp., and

0

DELTAX = (FINISH - START)/200.

RANGE      XMIN      XMAX

If the user has points in the data which fall outside the range of meaningful data, then this command will edit these data points from the fit; any point whose X value is below XMIN will not be included in the fit, and any point whose X value is above XMAX will also be omitted.

By default all points are included in the fit.

NORMATION

By default normation is set so that the constant "1" is multiplied by the function in THEORY.FOR. If the data is relative data, with no absolute significance, then issue the command NORMATION and the program will compute a normation factor for the fit. (ie. a constant multiplier other than "1").

FILE        [filename]

If the user would like to have a file created containing the extrapolated theoretical function and the parameters as determined by MINUIT, without all the other output produced by MINUIT, this command should be issued with an appropriate output filename. If no filename is included in the command line, the default filename will be MINUIT.DAT.

DEFSIG     [DEFSIG]

This command changes the default value of DY (ie. the default value of sigma) to suit the user. If this

## BIB KARL magnet spectrometer operating system HELP LIBRARY

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command is omitted the default assumed by the computer  
is DEFSIG=1.

Local commands are:

TITLE Title

A title for the data set will be included in the output.

SKIP

This command, when encountered before a data set, will  
cause the data set to be printed in the input, but the  
data set will NOT be included in the fit.

The data points are inputted as follows:

Xexperimental	Yexperimental	DYexperimental
"	"	"
etc.		

Notice that all the input is format free.

MINUIT

MINFIT

ERROR\_ANALYSIS

The errors that MINUIT calculates are the estimated errors of the current  
parameter values. These errors are directly related to the sigmas  
of the data points as provided by the user. For this reason, in order

to obtain meaningful error estimates on the parameters, it is necessary to provide meaningful error estimates on the input data. If you don't know the error in the input data, then MINFIT can produce estimates by itself, providing the function has been minimized once already with the original error estimates.

This redefinition of the sigmas of the input data is done by issuing the command

```
CALL FCN      6.
```

in the command file, or if you are working with a polynomial, the "error code" inputted to the MINFIT procedure should be 2.

Notice that if the sigma values are all equal, MINFIT computes the least squares fit for your data.

## MINUIT

### MINFIT

#### POLYNOMIAL

In many cases it will be desirable to fit the data to a simple polynomial of up to degree 5. For this case, a special option is available to the user that is particularly easy to use. In it, the polynomial is already defined, and the user need not worry about writing a theoretical function and producing a command file for MINUIT, the polynomial option of MINFIT does this for you.

The polynomial is of the form:

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$$F(X) = \text{Par}(1) + \text{Par}(2)*(X) + \text{Par}(3)*(X**2) + \text{Par}(4)*(X**3) + \dots$$

The parameters returned by MINUIT simply plug into this formula to provide you with the fitted function.

To run this option, simply produce an input file and follow the directions given to you by the MINFIT procedure which is executed with the command

\$ MINFIT

If the procedure is unable to provide you with a fitted function, try reducing the degree of the fitting polynomial.

MINUIT

MINFIT

THEORY

A sample THEORY function is given below:

```
CC*****
CC
CC
CC      FILE: THEORY.FOR      MINFIT MODULE THEORY
CC      =====            J.MEISSBURGER 3-NOV-82
CC                               MODIFIED: GREG SLAKOV 12.6.84
CC
CC*****
CC
CC
```

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Author: J.Meissburger

```
C  REAL*8 FUNCTION THEORY (PAR, NFLAG,XARG)
CC
CC  FUNCTION THEORY RETURNS THEORETICAL FUNCTION VALUE DEPENDING ON
CC  UP TO 55 PARAMETERS PAR(I)
CC
CC  NFLAG IS A SWITCH.  IT IS SET TO 1 FOR INITIALIZATION OF ALL OF THE
CC  CONSTANTS IN THE FUNCTION THEORY (IE., ANYTHING THAT DOES NOT DEPEND
CC  ON THE VECTOR PAR(I))
CC
C    IMPLICIT REAL*8 (A-H,O-Z)
CC
C  PARAMETER(MAX_POINTS=100,MAX_DSETS=20)
CC
C    COMMON /DATA1/ X(MAX_POINTS,MAX_DSETS),
C  *      Y(MAX_POINTS,MAX_DSETS),DY(MAX_POINTS,MAX_DSETS),
C  *      NSET,NDAT(MAX_DSETS),INPAR
CC
C    COMMON /FIT2/ I,II,COMF
CC
C  DIMENSION PAR(1)
CC
C  GOTO (100,200,300) , NFLAG
C100      CONTINUE
CC
CC      INITIALISATION STEP
CC      DEFINE (IE CALCULATE OR SET) ALL CONSTANTS OF THE FUNCTION
CC  IN THE SPACE BETWEEN LINE 100 AND RETURN.
CC
CC
C  RETURN
C200      XX=X(I,II)
CC
CC  THIS IS WHERE THEORY IS CALCULATED - DON'T WASTE ANY
CC  COMPUTING TIME DURING THIS STEP SINCE IT WILL LOOP
CC  THROUGH MANY TIMES
```

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```
CC
CC      DEFINE THE THEORETICAL FUNCTION STARTING AT LINE 250.
CC
C250      CONTINUE
CC
CC  THIS IS THE STANDARD POLYNOMIAL FUNCTION.
CC
CC  THEORY = PAR(1)+XX*(PAR(2)+XX*(PAR(3)+XX*(PAR(4)+XX*
CC      *      (PAR(5)+XX*PAR(6))))))
CC
CC      THIS FUNCTION IS A GAUSSIAN DISTRIBUTION, IT MAY BE
CC  USED TO FIT A GAUSSIAN PEAK
CC
C  THEORY = PAR(1)*EXP(-(((XX-PAR(2))**2)/PAR(3)))
CC
C  RETURN
C300      XX=XARG
CC
CC      THIS PART IS CALLED TO EVALUATE THE
CC      VALUE OF THEORY FOR THE EXTRAPOLATION POINTS.
CC
C  GOTO 250
C  END
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP MOCAL (14 July 1986):

MOCAL

\*\*\*\*\*

\*\*\*\*\* M O C A L \*\*\*\*\*

\*\*\*\*\*

Author: D.Paul

The Big Karl MORris chamber CALibration utility

Last update : January 1984

Most important commands : MOC> BKHELP COMMANDS  
To see an example : MOC> BKHELP EXAMPLE  
Standart run (one chamber plane) : MOC> BATCH GO  
Sign parameter in software : MOC> BATCH SIGN  
More commands : MOC> BKHELP MORE

=====

MOCAL

EXAMPLE

Typical procedure to get a new calibration for one plane:

- a) Type ACSTATUS and notify the actual version of ACON (filenames)  
Type ACPARAM to create the default (no parameter modification)  
version of ACON.
- b) Illuminate chamber by appropriate mistuning of Q2, Q3 and/or  
sweeping if necessary (white spectrum). Take list mode data  
on tape !



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c) Dump spectrum with ACON>DU,D and make a note in the logbook

d) EXIT ACON, stop subprocess and type MOCAL

```
MOC> READ ACAxxxx.SIN      ! read calibration spectra
MOC> ARRAY 1                ! define array # of wire spectrum
MOC> SCOPE                  ! display spectrum for check
MOC> FIND 300 1700          ! find peaks within limits of
                             ! wire number spectrum
MOC> DISPLAY FLUCT          ! check fluctuations (<4% !)
MOC> SCOPE 200 300          ! check beginning of spectrum
MOC> SCOPE 1500 300         ! check end of spectrum
MOC> KILL 1                 ! delete misidentified peak No. 1
MOC> STORE NX1 comment      ! store wire number calibration

MOC> ARRAY 2                ! specify array for drift time
                             ! analysis
MOC> SCOPE 1 2000           ! check spectrum
MOC> DRIFT 300 1500 16      ! get drift time spectrum binned
                             ! in 16 bins of same weight
MOC> DISPLAY WEIGHT          ! show result
MOC> STORE ND1              ! store drift time calibration

MOC> EXIT                   !
                             save calibration (Y/N) ! type Y to save calibration
```

e) Type ACPARAM and build ACON process including  
the new calibration by specifying

[USER.MORRIS]MOINI.FOR

as filename.

Hit <RET> when prompted for the calibration file

f) Hit <RET> when asked if you want to invoke MPS

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

>>>> G o o d   L u c k <<<< and watch what's  
coming out !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

MOCAL

COMMANDS

READ	read in calibration spectra ( single files)
ARRAY	get array to work on
FIND	find peaks in a wire spectrum and set windows
SWP	switch to a mode, where there is prompted for all param.
KILL	possibility for corrections
DRIFT	cut drift time spectrum into equal weighted bins
SCOPE	scope array - and , when calibration done: mark windows
DISPLAY	optical control on several parameters
STORE	store away done calibration of a parameter
BATCH	call batchfile ( ready programmed runs )
EXIT	save calibration (Y/N) Y

>>> More commands please type HELP MORE <<<

MOCAL

READ

READ filename [,from array to array]

filename	: name of a single-file	DEFAULT
array	: range of arraynumbers	read all arrays

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
>>> reads in arrays of a single file and stores them in a buffer
>>> use the command 'ARRAY' to restore an array from this buffer
>>> I F : exactly one arraynumber given THEN array is restored automatically
```

## MOCAL

### ARRAY

ARRAY      number

number                      : number of the array to be restored

```
>>> restore an array from the arraybuffer
>>> initialize for calibration on next parameter
```

## MOCAL

### FIND

FIND      lower channel,upper channel[,wire spacing,threshold,step,wobbel]

DEFAULT

lower channel      : approximate beginning of spectrum / should be given  
upper channel      : approximate end of spectrum                      when bad spectrum  
wire spacing      : approximate spacing between wire / given by program  
                    peaks  
threshold          : minimum counts in a peak  
                    to accept a peak  
step                : iteration step when searching peaks / 1 channel  
wobbel             : 1 = look for minimum between peaks / "1"

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Author: J.Meissburger

```
>>> find peaks in a wire spectrum and integrate them
}}}} in case of no success see also 'SWP'
```

MOCAL

DRIFT

DRIFT lower channel , upper channel [, number of bins ]

DEFAULT

lower channel : lower threshold for a valid event

upper channel : upper threshold for a valid event should be given

number of bins : number of equal weighted bins 16 bins

```
>>> cut drifttime spectrum into equal weighted bins
```

MOCAL

BATCH

BATCH [filename]

filename : filename or logical name for a batchfile

```
>>> call a ready programmed sequence of commands
```

```
>>> the program will prompt for parameter values
which are not given in the batchfile
```

```
>>> the logical name for the batchfile can be
defined before entering the program with the
command $ DEFINE ( VAX-VMS manuals )
```

# BIB KARL magnet spectrometer operating system HELP LIBRARY

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>>> logical names for the batchfiles  
[USER.MORRIS]GO.BAT,GOGO.BAT are  
automatically defined GO,GOGO when entering  
the program.

}}} example : MOC> BATCH GO  
calibration of one chamberplane

## MOCAL

### STORE

STORE      tablename ,    comment

tablename        : conventional name used for the calibration of  
                  the relevant parameter  
comment           : comment , which is written into the table header

>>> store away a ready done calibration  
}}} ( name-conventions should be looked up in the 'REFERENCE'-file )

## MOCAL

### SCOPE

SCOPE    lower channel [, range , compress factor ]

          DEFAULT

lower channel :                    first: scope all  
                                  then:escape from display  
range                : number of channels to    100ch or old value

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Author: J.Meissburger

be displayed  
compress factor : 'resolution' 1ch or old value

>>> scope current array

>>> the program is set to a mode , in which it replots  
after valid commands ( except DISPLAY ).

The command can be simply given instead of the  
lower channel ( e.g KILL , FIND , SET , FIX etc.)

>>> on return , the program exits the scopemode

>>> I F windows are present THEN they are marked.

The numbers are the peak or binnumbers.

See also: KILL

MOCAL

DISPLAY

DISPLAY option

LIMITS	: lower limits of windows ( default option)
FLUCTUATION	: fluctuation of spacings between window centroids - the difference from the mean value is given in percent
WEIGHT	: normated integrals of windows
CENTROID	: centroids of windows
WALK	: walk of limits compared to old calibration ( This needs a tablename as a parameter )

>>> x-axis = window number , y-axis = "option"

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

MOCAL

DISPLAY

WALK

WALK tablename

tablename : table of referencefile to be compared with  
new calibration

>>> compare current calibration with old calibration,  
the difference in channels is displayed.

MOCAL

CREATE

CREATE comment

>>> assemble new calibration file from stored tables ( see 'STORE' )  
>>> the reference file, which had been read in at the  
beginning of the program , is used as a pattern  
>>> reset all recalibrated tables , keep tables which are not  
recalibrated  
>>> dump this file on the default directory

MOCAL

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MORE

GET            read in a table of an old calibration file,  
              integrate current array with this calibration  
              (e.g. compute window integrals , centroids for display )  
RESTORE       restore table from memory,integrate current array  
              with this calibration.  
DIRECTORY     list names of stored tables  
TYPE [x]      type current table or stored table 'x'  
CLEAR         clear graphic display  
DUMP          dump for private use or when no reference file can  
              be given ( else use 'CREATE' or 'EXIT' )  
OFFSET        add offset on an old calibration table ( other  
              particles )

SET, FIX, ADD, SUBTRACT, HEADER for manipulating a table

MOCAL

CLEAR

>>> clear graphic display , nothing is initialized

MOCAL

DUMP

DUMP [filename,consistent routine,tables,mpstables,comment]



# BIB KARL magnet spectrometer operating system HELP LIBRARY

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filename : name of calibrationfile  
consistent  
routine : name of compatible fortran module (BK\_USER\_PAR)  
tables : these tables are dumped ( only last version )  
          DEFAULT: all tables stored  
mpstables : these tables will be converted to memphis  
          batchformat when using initialization module  
          "MOINI"

MOCAL

DIRECTORY

DIRECTORY

>>> list all stored tables, new versions already stored are marked with  
      "(recalibrated)"

MOCAL

TYPE

TYPE [tablename]

DEFAULT

tablename : name of stored table

current table

>>> type table on terminal

If no name is given type current table

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Author: J.Meissburger

MOCAL

INITIALIZE

INITIALIZE

>>> initialize windows, integrals etc. of the current table

MOCAL

SET

SET limit#,value [,limit#,value,...]

limit# : number of limit to be changed

value : value (channel)

values are allowed to be negative

MOCAL

ADD

ADD tablename

tablename : name of a stored table to be added

>>> perform a vektor addition of a table from the table directory

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and the current table

MOCAL

SUBSTRACT

SUBSTRACT      tablename

tablename : name of a table on the table directory to be  
             subtracted from the current table.

>>> see also ADD

MOCAL

FIX

FIX      lower limit , correction , upper limit , correction

lower limit    : first manipulated limit

correction     : value to correct lower limit

upper limit    : last manipulated limit

correction     : value to correct last limit

>>> generate a correction table .

All values between the lower and upper limit are computed  
by a linear interpolation between the corresponding corrections.  
This correctiontable is then added to the existing table.

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Author: J.Meissburger

MOCAL

HEADER

HEADER singlefile,arraynumber,size

DEFAULT

singlefile	: information in table header	old name
arraynumber	: information in table header	old value
size	: size of relevant parameter	old value

>>> size is important , as it might be a compiletime constant

MOCAL

SWP

SWP >>> switch between two modes

promptmode : prompt for all parameters ,  
Force prompted parameter input when using FIND

normalmode : short hand parameter prompting  
Restrict parameter prompting to important parameters

MOCAL

KILL

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

KILL peak# [,peak#,....]

peak# : peaknumber of peak to be deleted

>>> delete misidentified peak in wire spectrum .

}}} peaknumbers can be obtained from graphic display ( see SCOPE )

}}} KILL can be conveniently used in the 'scope-mode '.

the relevant part of spectrum is replotted

If peaks are missing type SWP

MOCAL

OFFSET

OFFSET value

value : offset to be added

>>> add constant offset on all nonzero elements of a  
calibration table

( correction of drifttimetable , eg. other particles )

MOCAL

GET

GET calibrationfile,tablename

DEFAULT

calibrationfile : old calibrationfile none

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Author: J.Meissburger

tablename : table to be read  
and restored in the  
buffer of the current  
array

>>> read in old calibration file for comparison  
see also RESTORE

MOCAL

RESTORE

RESTORE tablename

tablename : name of table to be restored

>>> restore table from table store (see DIRECTORY)  
>>> if an array is currently present ,the  
calibration will be applied on it :  
integrals & centroids of bins are computed .  
( see DISPLAY )

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

## BKHELP NEWS (14 July 1986):

### NEWS

```
*****
***** BK-VAX NEWS:  4-Jul-1986 *****
*****
```

News about the Big Karl operating system on VAX and PDP11

\*\*\*\*\*

To get a hardcopy use:

```
$ BKHelp/output=bkhelp.lis NEWS...    and
$ Print/delete bkhelp.lis
```

at DCL command level. The \$ BKHelp command provides you with the more detailed information on Big Karl operating system utilities. (See \$ BKHelp BKHELP)

### NEWS

### PRINTERS

Besides the standard systems line printer queue SYSS\$PRINT two more printers are available in the IKP main building, room 340 for either lineprinter-, memo- or letter-quality printing.

```
SYSS$LETTER      is queued to a LA100 letter/memo printer
SYSS$LASER       is queued to a LNO3 high quality laser printer
```

For more information see: \$ BKHelp PRINTERS...

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

NEWS

TERMINALs

For new terminal DCL and Fortran configuration support use:

\$ BKHelp TERMINALs... and

\$ BKHelp BKLIB BKUTY TT\*...

NEWS

SYSTEM\_layout

For information on the Big Karl VAX-11/780 site-specific system layout (in addition to VMS-HELP) use:

\$ BKHelp SYSTEM...

NEWS

RUNOFF

A sample RUNOFF input sample file is available to show most features used on the LNO3 laser printer. For more information use:

\$ BKHelp RUNOFF...



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP PATCHY (14 July 1986):

PATCHY

\*\*\*\*\* PATCHY on BK-VAX \*\*\*\*\*

\$ YPATCHY      CERN PATCHY main utility

-----

\$ YEDIT	Edit material of old files with cradle material to new files
\$ YINDEX	Print index for a PAM file
\$ YLIST	Card-for-card listing of a PAM file
\$ YSEARCH	Search a PAM file for cards containing any text string as specified by the cradle and possibly replace strings.
\$ YSHIFT	Copy OLD to NEW, activating/deactivating tagged cards and converting blank to single-C cards or vice versa.
\$ YCOMPAR	Compare NEW against OLD for identical contents.
\$ YTOBIN	Conversion card-image to compact binary
\$ YTOBCD	Conversion compact binary to card-image
\$ YTOCETA	Conversion from binary to CETA ( tape must be mounted /FOR )
\$ YFRCETA	Conversion from CETA to binary ( tape must be mounted /FOR )

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP PLSORT (14 July 1986):

PLSORT

\*\*\*\*\*

\*\*\* P L S O R T \*\*\*

\*\*\*\*\*

Author: L.Zemlo, C.Sukoszd

VERSION 12-Dec-84

MEMPHIS and ND6600 general purpose sorting utility

Sorting utility for BIGKARL and ND6600 list mode data providing user-defined pseudo-ADC's and parameters.

When you execute PLSORT at DCL command level you will first be asked for a Fortran file specifying BK\_USER\_INI

This subroutine will be linked and executed as task PLINI to do initial parameter and table value computations. It will also define the filename of a subroutine BK\_USER\_PAR to redefine the pseudo-ADC's MPS\_ADC(1 .. 16) .

If your don't need pseudo-ADC's just type <RET> for the filename specifying BK\_USER\_INI.

To define extra user parameters PARAM(1 .. 30) you will then be asked to specify the name of a file providing a subroutine DO\_PARAMETERS. Typing <RET> will default all parameters to be 2K copies of the MPS\_ADC's.

The header of your DO\_PARAMETERS definition file (separated by a form feed <FF> from the program source text) will be copied to the file PLSORT.PAR and used by PLSORT for documentation.

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

## PLSORT

### Input

List mode data from MPS (MEMPHIS system) or MPA (ND 6600 system).

Decoded data are filling arrays MPS\_ADC (INTEGER\*2 type) in  
COMMON /BK\_EVENT/, see BKCOMM.FOR

## PLSORT

### Pseudoparameters

Pseudoparameters are defined (max number 30) by user in DO\_PARAMETERS  
subroutine (array PARAM (INTEGER\*4 type) in COMMON/PARCOM/).

Their values should lay in range 1 - PAR\_SIZE.

PAR\_SIZE must be defined for each pseudoparameter in the same subroutine  
(max PAR\_SIZE = 2048).

The values of pseudoparameters outside given range are neglected.

## PLSORT

### Gates

The user can define up to 40 gates in coincidence or anticoincidence.  
Among them up to 10 can be defined as 2-dim gates (areas of interest)

## PLSORT

### Complex gates

User can define up to 10 complex gates in coincidence or anticoincidence. The complex gates from 81,83,85...99 are in coincidence, the gates 82,84,...100 are in anticoincidence. For the condition list of these complex gates every preceding gate number can be used (also preceding complex gate numbers!). For more information see CONLIST and DCOM !

## PLSORT

### Arrays

User can define up to 30 1-dim arrays (size automatically defined by size of chosen pseudoparameter) and max two 2-dim arrays .

The 2-dim arrays may have low resolution (128\*128 points), or high resolution (640\*640 points). For high resolution plots there is no possibility of making a projection (see. CUT ).

Filling of those arrays can be conditioned by up to 10 conditions chosen from previously defined gates. For more information see ARRAY, CONLIST and SCOPE 2DIM.

## PLSORT

### Units

## BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

PLSORT uses the following input/output units:

TTY as input (interactiv mode) - unit number 5

TTY as output (interactive mode) - unit number 6

Temporary unit used for all transfers of data - unit number 30

Procedures batch-files as input - unit numbers 41,42,43,44....

Events (list mode data) - unit number 32

PLSORT.LOG (full documentation of program execution) - unit number 33

### PLSORT

#### Constants

Up to 512 constants stored in CNST array of COMMON/CON/ can be used for defining pseudoparameters in DO\_PARAMETERS subroutine. First 20 of them are equivalenced with constants C1,C2, ... ,C20.

### PLSORT

#### MESSAGE

Message line starts with '!' or ';' after prompt PLS>.

Copied to PLSORT.LOG file without affecting the execution of the program.

### PLSORT

#### ADD

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Add two 1-dim arrays: ARRAY1 + ARRAY2 and locate product in ARRAY3

Transfer of channels according to the formula:

$$a1*ARRAY1+a2*ARRAY2+offset \Rightarrow ARRAY3$$

Format:

AD[D] i1,i2,i3,a1,a2,offset<CR>

Comment string (one line,not more then 64 characters)<CR>

Default

i1...3 - number of ARRAY1...3 (value 1-30) none

a1,a2 - values of multiplication factor 1.

offset - offset 0.

PLSORT

BATCH

Terminates execution of a Procedure batch-file. This command `m u s t` be the last statement in each procedure-file (see example below).

Procedure files can be called using the PRO command (see there).

Exeptionally for calling procedure-files from the interactive level the BATCH command is also allowed (for keeping consistency with previous convention).

Format: B[ATCH] [filename]

filename must be given only for calling a procedure-file from the interactive level !

Procedure file structure:

PLSORT INPUT DATA <-- first line

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
.  
.  
      <-- input commands  
.  
BATCH      <-- last line
```

## PLSORT

### PROCEDURE

Format: PR filename

You may build procedures from sequences of valid PLSORT commands.  
These commands will be executed by calling the file which contains them.  
The PR[OCEDURE] command is used to call a procedure file.  
Procedure files can be nested the same way as subroutines in Fortran.  
(For calling a procedure file from the interactive level you may  
use - exceptionally - also the BAT command <see there> ).

Procedure file structure:

```
PLSORT INPUT DATA  <--- this must be the first statement  
.  
.  
      <--- These are valid PLSORT commands (Including PRO !!)  
.  
.  
BAT      <--- this must be the last statement
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

PLSORT

CHANGE

Change channel setting in gate definition

Format: CH[ANGE] ng,np,nl,ng

	Default
ng - gate number	none
(if ng<0 then the last ABS(ng) gate-pairs will be deleted!)	
np - parameter number	old value
nl - lower channel	old value
nu - upper channel	old value

PLSORT

REPLACE

Replace definition of 1-dim array

Format: REP[LACE] na,np,con\_list

na - array number

np - parameter number

con\_list - list of gates (till 10) terminated by 0  
(for more details see CONLIST)



. BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

PLSORT

CLEAR

Clear (zero) all defined arrays

Format: CL[EAR]

Attention: Arrays which were not explicitly defined  
are not erased ! (e.g. saved arrays)

PLSORT

SET

Set values of constants in COMMONS: CON,INF,STR

PLSORT

SET

CON

Set value for constant used in user DO\_PARAMETERS subroutine

Format: SE[T] C[ON] n,v

Default

n - number of element in CNST array · none

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

v - value of CNST(N)                      old value

PLSORT

SET

INF

Used to change actual configuration of program  
(see listing of PLCOMM.FOR)

Format: SE[T] I[NF] n,iv-

Default

n - number of element in COMMON /INF/    none

iv - value of this element                  old value

Don't use if you are not absolutely sure what are you doing  
- you can make terrible mess !!!

PLSORT

SET

STR

Change string in COMMON /STR/ or restore array

These strings are used to create standard filenames:

String # 4 - filename of list mode data from MEMPHIS  
(eg. MTAO:ACA\$\$\$\$.LMO)

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

- # 5 - filename for dump file
- # 6 - filename for definition file
- # 7 - filename for BATCH input file
- # 8 - filename for SHOWPLOT file

Format: SE[T] S[TR] n,new string<CR>

n - number of string replaced by new string  
if n<0 then restore(activate) array -n

## PLSORT

### CUT

Project part of 2-dim array on X or Y axis

Result saved in array # 30

Format: CU[T] istart,data1, ... ,dataN,0  
where dataI consists of 4 values: ax,ia,nl,nu

	Default
istart - define location of cut spectrum in array # 30 (beginning channel = istart*128+1)	0
ax - Y or X for projection on X or Y axis, respectively	none
ia - 2-dim array number (31 or 32)	31
nl - lower channel of cut in ax axis	1
nu - upper channel of cut in ax axis	128

THERE IS NO POSSIBILITY OF PROJECTION FOR HIGH RESOLUTION PLOTS !

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

PLSORT

2DGATE

Define 2-dim gate (area of interest)

Format: 2DG[ATE] ng,nc,npx,npj

Where: ng - gate number (from 79 downto 61)  
(if ng<0 then last ABS(ng) 2DG-pairs will be deleted)  
nc - contour number (must be previously defined)  
npx,npj - parameters in X and Y axis respectively

PLSORT

CONTOUR

Define contour for 2-dim gate (area of interest)

Format: CO[NTOUR] list\_of\_coordinates

Where list\_of\_coordinates - set of pairs of X,Y coordinates separated  
by comma or space and terminated by <CR>

PLSORT

DCOM

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Define complex gates:

User may define up to 10 complex gates (gate numbers 81 to 100). The complex gates with odd number are in coincidence, the ones with even number are in anticoincidence.

These complex gates can be used for conditioning one- or twodimensional arrays during the sort the same way, as the simple gates.

Format: DC[OM] gn conlist

Where

gn is the complex gate number to define. Default: next free  
(If  $gn < 0$  then last  $ABS(gn)$  complex gate-pairs are deleted)  
conlist List of gate numbers (maximum 10). Default: none  
(Terminated by 0)  
In the condition list every existing gate number  
may appear, which are less than gn.  
For more information see CONLIST.

PLSORT

DELETE

Delete 1-dim array from list of arrays

Format: DEL[ETE] n

Where n - array number

If  $n < 0$  then last  $ABS(n)$  arrays are deleted.

PLSORT

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

DFL

Set default values for configuration definition

Format: DF[L]

Function:

- 1) defines 10 1-dim arrays for first 10 parameter (without condition)
- 2) releases all gates and contours
- 3) deletes arrays 11-32
- 4) clears CNST array in COMMON /CON/
- 5) sets default values for strings in COMMON /STR/
  - # 4 - MTAO:ACA\$\$\$\$.LMO
  - # 5 - PLAS\$\$\$\$.DMP
  - # 6 - PLAS\$\$\$\$.DEF
  - # 7 - PLAS\$\$\$\$.BAT
  - # 8 - PLAS\$\$\$\$.PLO
- 6) sets default values for WINDOWS

PLSORT

SCOPE

Display 1-dim or 2-dim array !!! Illegal in batch mode !!!

PLSORT

SCOPE

1-dim

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Display 1-dim array

Format: SC[OPE] na,nl,nu,y\_max

Default

Where: na - array number (1-30)	none
nl - beginnig channel of display	old value for this array
nu - end channel of display	old value for this array
y_max - maximal value of vertical axis	0

If y\_max=0 then maximal value of vertical axis is equal to maximum number of counts per channel in displayed range (but y\_max>9)

PLSORT

SCOPE

1-dim

Array

Enables you to plot another array between the previously defined channels but with a new scale factor WITHOUT ERASING the display. So you can plot a spectrum over another one and can directly compare them. Always the parameters of the LAST spectrum are kept ! Execution: Cursor in displayed frame, then press A and answer the questions. Default values are always the last ones.

PLSORT

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

SCOPE

1-dim

Clear

Clear displayed range of array.

When range 1-2048 is displayed make array nonactive

Execution - CURSOR in displayed frame and press C

PLSORT

SCOPE

1-dim

Expand

Expand part of display between markers

Execution - CURSOR between markers and press E

PLSORT

SCOPE

1-dim



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Full

Display array between channels 1-2048

Execution - CURSOR in displayed frame and press F

PLSORT

SCOPE

1-dim

Integrate

Integrate part of the array between markers

Execution - CURSOR between markers and press I

PLSORT

SCOPE

1-dim

Logarit

Log scale in Y-axis

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Execution - CURSOR in displayed frame and press L

PLSORT

SCOPE

1-dim

Mark

Set marker

Execution: Move CURSOR to chosen channel and press M

On display returned channel number and number of counts in this channel

PLSORT

SCOPE

1-dim

Outlp

Write in \*.LOG file contents of array between markers

Execution - CURSOR between markers and press 0

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

PLSORT

SCOPE

1-dim

Point

Change display from histogram to point

Execution - CURSOR in displayed frame and press P

PLSORT

SCOPE

1-dim

Quit

Return to PLS>

Execution - CURSOR in displayed frame and press Q

PLSORT

SCOPE

1-dim

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Replot

Return to histogram mode and replot display

Execution - CURSOR in displayed frame and press R

PLSORT

SCOPE

1-dim

Scale

Replot with changed Y-scale

Execution: Press S and type new y\_max value

PLSORT

SCOPE

1-dim

Window

Store information about range and all markers of displayed arrayes.  
These information may be stored in 9 registers called back

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

by pressing numbers 1-9.

Execution: Press W and type register number.

Default definition: range markers

# 1	1-256	1,128,129,256
# 2	257-512	257,384,385,512
# 3	513-768	513,640,641,768
# 4	769-1024	769,896,897,1024
# 5	1025-1280	1025,1152,1153,1280
# 6	1281-1536	1281,1408,1409,1536
# 7	1537-1792	1537,1664,1665,1792
# 8	1793-2048	1793,1920,1921,2048
# 9	1-2048	1,128,129,256,257,384,385,512,513,640,641,768, 769,896,897,1024,1024,1152,1153,1280, 1281,1408,1409,1536,1537,1664,1665, 1792,1793,1920,1921,2048

PLSORT

SCOPE

2-dim

Display 2-dim array

Format: SC[OPE] na,min,max

Where:

Default

na - array number (31 or 32)

31

min - lower level in number of counts

1

max - upper level in number of counts

1000000

Function: Displays coordinates corresponding to channels containing

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

number of counts between min and max.

For high resolution plots the 640\*640 channels are grouped in 128\*128 groups, containing 5\*5 channels each. The content of these groups is incremented during the sort, though it is maximized to 127. During the display procedure, only those groups are (or are not) plotted, whose content is between min and max.

PLSORT

SCOPE

2-dim

Cursor

Activate cursor

Execution: Press C, to get channel coordinates move cursor to chosen place and press space. You get coordinates of nearest channel.

PLSORT

SCOPE

2-dim

Define contour

Define contour for 2-dim gate

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Execution: Press D, then enter the contour number to define (if you enter -1, the last contour will be deleted). Then move cursor to the chosen position and press D to store the coordinates. For finishing contour definition press E.

PLSORT

SCOPE

2-dim

Replot

Replot with new values of min and max

Execution: Press R and give required new values of min and max

PLSORT

SCOPE

2-dim

Quit

Return to PLS>

Execution: Press Q

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

PLSORT

SCOPE

2-dim

Show contour

Show previously defined contour

Execution: Press S and type contour number

PLSORT

SCOPE

2-dim

End

Termination of contour definition

Execution: Press E

PLSORT

DUMP



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Save or read saved configuration (PLSORT internal format)

Format: DU[MP] key,filename

key - keyword to control DUMP function

filename - name of file of saved configuration

List of keywords:

R - read full configuration

RALT - read full configuration of sorts before FEB-1983

RINF - read configuration definition only

W - save full configuration - this keyword does not want filename

WD -save full configuration - this keyword wants a filename

(If the destination file already exists, it asks for replace.

In batch mode this option does not replace the existing file,  
but leaves the DUMP subroutine.)

WINF - save configuration definition only

WSIN - write arrays in format compatible with VAXUNF program

PLSORT

EXIT

Exit program and close all files

Format: E[XIT] Y[ES]

PLSORT

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

GATES

Define 1-dim gate

Format: G[ATE] np,nl,nu

Where: np - parameter number

nl - lower limit

nu - upper limit

PLSORT

INTEGRATE

Integrate between channels incl. background subtraction

Format: I[NTEGRATE] na,nl,nu

Where:

Default

na - array number (1-30)

none

nl - lower channel

1

nu - upper channel

last channel of array

PLSORT

NDSORT

Sorting of MPA list mode data - Nuclear Data 6600

Before calling PLSORT tape must be mounted: MOU MT.../FOR

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

## Format:

ND[**SORT**] ntu,filename,nb,nr ---- if ntu= 0 or 1 (Tape Sort option)  
2,S,filename1,nb,nr ---- Disk Sort option  
2,C,filename1,ntu,filename - Copy from tape to disk

## Where:

## Default

ntu - number of I/O unit (0 or 1 for tape, 2 for disk )	
filename - name of file with data on tape	none
filename1- name of I/O file on disk	none
nb - number of first record to be sorted (nb>3)	actual position
nr - number of records to be sorted	999999

## PLSORT

## PARAMETERS

Display and print present configuration and check set-up.

Format: PA[**RAMETER**] key

key - optional: if key(1:1)= P, print configuration on line printer  
if key = [P]E print configuration errors  
if key = [P]W print warnings and errors  
if key = [P]C print cross-references, warnings and errors

## PLSORT

## READ

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Read one 1-dim array form DUMP file or from SHOWPLOT-format file

Format: RE[AD] filename,ns,nd

Where: filename - name of DUMP or SHOWPLOT file (source)

ns - array number in source file

ns must be -1, if transferring a SHOWPLOT file !!!

nd - destination array number

PLSORT

SAVE

Manipulate (expand, compress, shift) and save 1-dim array

Format: SA[VE] ns,nd,shift,expand<CR>  
comment<CR>

Where:

Default

ns - source array number

none

nd - destination array number

none

shift - shift

0.

expand - expand (or compress when <1 )

1.

comment - one line but not more then 64 characters

PLSORT

ARRAY

Define 1-dim array (spectrum) and conditions

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Format: AR[RAY] np,con\_list

Where: np - parameter number

con\_list - list of previously defined gates (max=10)

(for more details see CONLIST)

PLSORT

CONLIST

There is a possibility of forming logical conditions using the previously defined one or two-dimensional gates. The following example explains the use of them.

Let A,B,C and D be gate numbers. The following condition-list

A B D -B C -A D

means:

$(A \text{ AND } B \text{ AND } D) \text{ OR } (B \text{ AND } C) \text{ OR } (A \text{ AND } D)$

Up to 10 items may appear in a condition list. Naturally, anticoincidence gates may also be used. One- or two-dimensional arrays can be conditioned during the sort with the condition-lists.

PLSORT

SORT

Sort MEMPHIS list mode data

When sorting from tape, tape must be mounted before calling PLSORT program !

BIB KARL magnet spectrometer operating system.HELP LIBRARY

Author: J.Meissburger

Format: SO[RT] filename,nb,nr,rew\_key

Where:	Default
filename - name of file with data	none
nb - number of first record to be sorted	actual position
nr - number of records to be sorted	999999
rew_key - if 'Y' tape will be rewinded before searching for file	

!!! rewkey is read only when sort is from magtape !!!

PLSORT

2DARRAY

Define new or change one of two previously defined 2-dim arrays

(See also by SCOPE 2DIM)

Maximum number of 2-dim arrays = 2

Format: 2DA[RRAY] (key,na,) nhr,npx,nox,ncx,npj,noy,ncy,con\_list

Where:	Default
(key,na) - read only when array # 32 is already defined if key='YES' then array # na is redefined else nothing done	
nhr - For nhr=0 ---> Low resolution, else high res.	0
npx - parameter in x axis	none
nox - offset in x axis	0
ncx - compression factor in x axis	16
npj - parameter in y axis	none
noy - offset in y axis	0
ncy - compression factor in y axis	16
con_list - list of previously defined gates (max=10) terminated by 0<CR>	

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

(for more details see CONLIST)

PLSORT

WRITE

Write 1-dim spectrum into file compatible with SHOWPLOT program

Format: W[RITE] filename,na<CR>  
comment<CR>

Where:

filename - destination file

na - source array number

comment - comment (max 80 characters)

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP PRINTERS (14 July 1986):

PRINTERS

Three line printers queues are available on the Big Karl VAX,

SYS\$PRINT	COT-Printronix system line printer
SYS\$LETTER	LA-100 memo/letter quality printer
SYS\$LASER	LN03 high quality letter printer

Formatting is supported by a set of print forms, see:

\$ show queue/form,

by logical configuration files for device and RUNOFF formatting  
(e.g. LASER\$CONFIG\_ELITE), and by DCL command procedures (e.g. \$ LASER  
or \$ LPRINT)

PRINTERS

Printronix

The Printronix lineprinter is the standard systems line printer connected to the line printer queue SYS\$PRINT. The only special support on this printer is the Calcomp emulator library BKLIB:CALTRON.OLB which is used by some of the Big Karl utilities.

PRINTERS



## LN03

The LN03 laser printer is supported by the following print command procedures:

\$ LASER filename [options]	for command formatting
\$ MPRINT filename	for 132 char/line printing
\$ NPRINT filename	for internally formatted letter printing

For standard (german character set) letter printing the combined RUNOFF and device configuration files:

laser\$config_elite	(12 pt. Elite) or
laser\$config_courier	(10 pt Courir)

may be requested within a RUNOFF input file (\*.RNO) and formatted with

\$ RUNOF filename (LN03 device specific RUNOFF)

and printed on the laser printer with

\$ NPRINT filename.LNI

Parameters to the DCL command "LASER" are explained on the header page produced by using the command itself !

More logical configuration files are available, see:

\$ show logical laser\$\*

Files translating to \*.DAT are device-only configuration, \*.RNO are RUNOFF-only or combined configuration files.

## PRINTERS

### LA100

The LA100 correspondence printer is supported by the command:

```
LPRINT filename [U or G] [L or S]    for
```

U = ASCII character set

G = ISO german character set

L = large (letter) print format and quality

S = small (lineprinter) format and quality

If you omit the filename the command will prompt you for additional parameters. Typing <RET> for the filename allows you to configure the letter printer for a desired character set and print quality to print files containing no internal configuration data.

If you explicitly use any of the letter\$... configuration files in your text (See: BKHelp RUNOFF) this will override the LPRINT parameter specifications !

The most important logical configuration file (combined device and RUNOFF configuration) is

```
letter$config
```

which configures the LA100 for german character set, 12 pt. letter gothic, high quality printing. For more configuration files see:

```
$ show logical letter$*
```

BKHELP RDSHOW (14 July 1986):

RDSHOW

RDSHOW is a template program to show the use of the Big Karl standard free style input routines. To get a source listing see:

\$ BKHelp rdshow source

General:

The RD... routines allow format-free terminal input with a set of interesting features that are intrinsic to the routines:

Free style command parsing with automatic defaulting  
Forcing of variable types with lookahead for re-interpretation  
Automatic background logging mode of terminal input  
Program level batch mode with or without echo  
Automatic Journal/Recovery mode for session reply  
(This may be used to easely split a program with interactive input to calculate in a batch queue while maintaining the interactive input without reprogramming. For details ask J.Meissburger )

RDSHOW

source\_code

CC

CC

CC     F I L E :     RDSHOW.FOR

CC     =====

CC

SAMPLE PROGRAM TO DEMONSTRATE THE  
STRUCTURE OF A PROGRAM USING THE  
FREE STYLE INPUT UTILITIES RD...

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC                                     J.MEISSBURGER 20-AUG-82
CC
CC *****
CC
CC THIS FILE SHOWS AN EXAMPLE OF A COMMAND PARSER USING THE FREE
CC STYLE INPUT ROUTINES RD...
CC
CC THE SUBROUTINE "SERVICE" IS A DUMMY ROUTINE TO BE REPLACED BY
CC APPROPRIATE SERVICE SUBROUTINES. THE RETURN FLAG "LFLAG" HAS
CC TO BE RETURNED +1 ON SUCCESS, 0 ON ERROR HANDLED WITHIN THE
CC SERVICE SUBROUTINE AND -1 ON UNRECOGNIZED COMMAND ERROR.
CC
CC ALL ROUTINES WITH A FEW EXCEPTIONS AS INDICATED ARE WRITTEN IN
CC STANDARD PDP11 FORTRAN AND SHOULD RUN ON ANY COMPUTER. (REPLACE
CC "BYTE" BY "LOGICAL*1" IF NOT SUPPORTED!)
CC
CC *****
CC Note: RD_HELP is specific to VMS systems, all other routines can
CC          run on any computer providing at least Fortran IV
CC *****
CC
CC
CC      PROGRAM RDSHOW
CC
CC      INTEGER*4 CMDS(4), CMD, NC, LFLAG
CC      INTEGER*4 LCMDS(4)
CC      CHARACTER*128 HELP_LIBRARY, LEVEL_1_KEY
CC
CC LIST OF VALID COMMANDS
CC
CC      DATA NCMS /4/
CC      DATA CMDS /'EX','HELP','CD3','CD4'/
CC      DATA LCMDS /2,4,3,3/
CC
CC
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC INITILISE TERMINAL I/O, PROGRAM DEFAULTS AND DATA FILES
CC AND PRINT PROGRAM IDENTIFICATION.
CC
C    CALL INIT (LFLAG)
CC
CC HERE TO READ NEXT COMMAND LINE
CC
C10  CALL RDCMD(CMDS,LCMDS,4,'RDC',CMD,NC,LFLAG)
CC
CC CHECK FOR EOF DURING COMMAND INPUT
CC
C    IF (NC.EQ.0) GOTO 100
CC
CC BRANCH TO COMMAND SERVICE
CC
C    GOTO (100,200,300,400) , NC
CC
CC EXIT , THIS COMMAND PROVIDES STANDARD EXIT TO DCL
CC
C100 CALL SERVICE (NC, CMD, LFLAG)
CC  CALL CLEANUP ( ..... )
C    CALL EXIT
CC
CC HELP
CC
CC  This routine uses system services provided by VAX-11 VMS
CC  to gain keyed access to the HELP library BKLIB:BIGKARL.HLB
CC  To include a level-1 HELP file in the Big Karl system help
CC  library please inform J.Meissburger
CC
C200 HELP_LIBRARY = 'BKLIB:BIGKARL'
C    LEVEL_1_KEY = 'RDSHOW'
C    CALL RD_HELP ( HELP_LIBRARY, LEVEL_1_KEY )
C    GOTO 10
CC
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC CD3
CC
C300  CALL SERVICE (NC, CMD, LFLAG)
C      GOTO 10
CC
CC CD4
CC
C400  CALL SERVICE (NC, CMD, LFLAG)
C      GOTO 10
CC
CC
C      END
CC
CC *****
CC
CC
C      SUBROUTINE INIT (LFLAG)
CC
CC INITIALISE PROGRAM
CC
C      COMMON /INOUT/ NIN, NOUT
CC
C      DIMENSION DAY(3), TIM(2)
CC
CC INITIALISE TERMINAL I/O ON LUNS 5/6 IN COMMON /INOUT/
CC
C      CALL RDSET (0, 5, 6)
CC
C      CALL DATE(DAY)
C      CALL TIME(TIM)
CC
C      WRITE (NOUT,20000) DAY, TIM
C20000      FORMAT(// ' RDSHOW: ',3A4,2X,2A4/)
CC
CC LFLAG = 2 FOR THE VERY FIRST CALL TO RDCMD !
```

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
CC
C      LFLAG = 2
C      RETURN
CC
C      END
CC
CC *****
CC
CC
C      SUBROUTINE SERVICE (NC, CMD, LFLAG)
CC
C      INTEGER*4 NC, CMD, LFLAG
CC
C      COMMON /INOUT/ NIN, NOUT
CC
C      WRITE (NOUT,20000) NC,CMD
C20000      FORMAT (' SERVICE # ',I2,'      CMD = ',A4)
C      LFLAG=1
C      RETURN
CC
C      END
```

RDSHOW

CD3

..... Dummy command CD3 of RDSHOW executed ! .....

RDSHOW

CD4

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Dummy command CD4 of RDSHOW executed !

RDSHOW

HELP

Access HELP information of RDSHOW in Big Karl system  
help library.

RDSHOW

EXIT

Exit task and return control to DCL



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP RSX11M (14 July 1986):

RSX11M

PDP-11/40 and RSX-11M:

-----

Login as:        username: PDP\_USER    (turnkey account)  
                 password: USER

or                username: PDP\_familyname resp.

to run full RSX-11M compatibility mode (UIC=[100,100]). Task images will have extension .EXE instead of .TSK, and the TKB command files have to include ..... , LB:[1,1]FOROTS/LB, in order to link in the complete RSX run time library.

To test CAMAC software on the VAX the CAMac SIMulation Inter-face Routines corresponding to the IKP-Library on the PDP-11/40 (Big Karl) are available as .... , DK3:[1,4]CASIMIR/LB, .....

The Fortran compiler, linker, macro and librarian are identical to those running on the Big Karl PDP-11/40 control computer ! Device-dependent system routines have nevertheless to be linked in on the PDP11 in order to work properly (There is no translation of hardware addresses and interrupt vectors from the PDP11 to the MCR-compatibility mode)

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP RUNOFF (14 July 1986):

RUNOFF

To support high quality text printing (large format) use:

```
$ RUNOF filename ( n o t  RUNOFF ) for the LN03 laserprinter
or
$ RUNOFF filename                for the LA100 printer
```

There are several combined Runoff/device-configuration files available for text formatting and font selection. The most important ones are:

```
laser$config_elite      for elite 12 pt (german) printing
laser$config_courir     for courir 10 pt (german) printing
```

on the LN03 laser printer, and

```
letter$config           for letter-gothic 12 (german) printing
```

on the LA100 letter printer.

You may include the configuration files by a Runoff request:

```
.req "letter$config"    for combined RUNOFF and letter printer
                        configuration (high quality letter
                        gothic, german)
```

(which is the best method since it automatically includes possible configuration file updates), or you may copy the configuration data directly into the text file by a EDT "Include" command.

In case you wish a different layout for the LA100 letterprinter you should use:

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

```
.req "letter$rho"      default RUNOFF configuration
.lo xx,xx              to define your layout
.req "letter$default"  for letter printer setup
```

as first commands in your RNO file. The layout for laser printer configuration files is undefined , use of .layout 3,2 as standard is recommended.

The commands for printing are:

```
$ LPRINT filename.MEM      for letter printer output   or
$ NPRINT filename.LNI      for laser printer output
```

For more information also see: \$BKHelp PRINTERS

## RUNOFF

### LN03\_Example

To see a sample input file using most of the character- and font-switching features of Runoff and the laser printer you may look into the sample file:

```
$ edt/read user$manager:lnetest.rno
```

or produce a hardcopy output with:

```
$ runof user$manager:lnetest
$ nprint/delete lnetest.lni
```

This sample output shows USASCII, German, DEC-supplemental character sets and examples of general text formatting including scientific formulae.



Zusammenstellung der Daten über erreichbare Teilchenzahlen  
bei der Stripping-Injektion von  $H_2^+$  in COSY

Die Injektion von Protonen in COSY wird über die Stripping-Reaktion von  $H_2^+ \rightarrow 2$  Protonen erfolgen. Die Wahrscheinlichkeit für die Umladungsreaktion beträgt bei einem  $20 \mu\text{g}/\text{cm}^2$  dicken Kohlenstofftarget ca. 90%. Daher beziehen sich die nachfolgenden Berechnungen der erreichbaren Protonenzahlen auf ein Kohlenstofftarget dieser Dicke.

Injektionspunkt ist die Austrittsöffnung am Dipolmagnet MD23 (der vorletzte Dipol der Einheitszelle vor dem Targetteleskop, Entfernung vom TP1 in Umlaufrichtung 158.147 m).

Bei der folgenden Berechnung der erreichbaren Anzahl der Protonen im Ring gehe ich von einem zur Verfügung stehendem  $H_2^+$ -Strom von  $5 \mu\text{A}$  aus. Dies ergibt einen Protonenstrom von  $I_p = 10 \mu\text{A}$ .

Dies entspricht einer Protonenanzahl pro sec.  $P_0$  von:

$$P_0 = \frac{I_p}{1.6 * 10^{-19} \text{ A sec}} = 6.25 * 10^{13} \text{ Protonen/sec}$$

$$= 1.25 * 10^8 \text{ Protonen pro } 2 \mu\text{sec}$$

die zeitlich einem Umlauf der Protonen in COSY entsprechen.

Die angenommene  $H_2^+$ -Energie beträgt 90 MeV, entsprechend einer Protonenenergie von  $E_p = 45 \text{ MeV}$ .

Die Berechnungen beziehen sich auf den Arbeitspunkt, der im technischen Bericht in der Tabelle 3 als Arbeitspunkt für die Injektion beschrieben ist.

Parameter sind:

Arbeitspunkt :  $Q_x = 3.867$        $Q_z = 4.119$

Lattice-Funktionen am Ort der Injektion:

$\beta_x = 7.031 \text{ m}$        $\beta_z = 15.426 \text{ m}$   
 $D = 4.364 \text{ m}$

Maxima in den Einheitszellen:

Lattice-Funktionen :  $\beta_x = 25.080 \text{ m}$        $\beta_z = 15.913 \text{ m}$   
Dispersion :  $D = 7.603 \text{ m}$

Die horizontale Akzeptanz ist bestimmt durch den Ort der maximalen Betafunktion und maximalen Dispersion. Die Maxima beider Funktionen treten in dem Kühlteleskop mit den Werten  $\beta_x = 46.91 \text{ m}$  und  $D_{\max} = 10.2 \text{ m}$ . Bei einer möglichen Strahlausdehnung von  $d = \pm 65 \text{ mm}$  und einer Impulsunschärfe von  $\delta = \pm 0.2 \%$  ergibt dies eine monochromatische Akzeptanz von:

$$\epsilon_x = \frac{(65 \text{ mm})^2 - (101.2 \text{ mm}/\%)^2 * (0.2 \%)^2}{46.91 \text{ m}} = 81 \text{ mm mrad}$$

Sample formula:

$$\frac{\dot{\Phi}^2}{2} - \frac{\Omega^2}{\cos \Phi_s} (\cos \Phi + \Phi \sin \Phi_s) - \frac{2 \pi \Omega^2}{eV_0 \cos \Phi_s} E_1 \Phi = C$$

The input text file is available as USER\$MANAGER:LNTEST.RNO  
and may be inspected with: \$ EDT/READ USER\$MANAGER:LNTEST.RNO



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP SYSTEM (14 July 1986):

System\_layout

The Big Karl site-specific system layout conforms to required VMS standards identifying every user uniquely on the system.

Apart from the user directories there are three main directory roots to store the DEC-VMS and Big Karl magnet spectrometer operating system:

sys\$sysroot: DEC-VMS system root (not site-specific)

user\$sysroot: Big Karl site specific system components:

user\$startup: System startup files

user\$manager: Device and system support

user\$login: Login/logout procedures

user\$system: System wide (installed) images

userdisk:[bksys...] Big Karl operating system  
application software (This root  
keeps the reference copies of all  
operating system components and  
should be maintained with great care !)

For each IKP-group (such as BK, KR etc.) there is a system wide logical group device ( BK\$: ) and a corresponding master group directory ( BK\$LOGIN: ). Login USERNAME is the group identifier ( BK ) followed by the family name of the group member ( BK\_MEISSBURGER ) separated by an underscore "\_". All group members have unique UIC member numbers but share the UIC group number with the master UIC.

Groups supported so far are:



## BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BK	Big Karl magnet spectrometer
C	C O S Y
CYC	Cyclotron
KR	Kernreaktionen
AXN	(alpha, Xn)-group
PKL	Peter Kleinheinz group
JOS	Josef group

Symbol definitions relevant to a l l group members should be entered into the group LOGIN file ( e.g. BK\$LOGIN:LOGIN.COM ) preferably by a group manager who is responsible for a group-wide systems management (username = group identifier followed by "\_", e.g. "BK\_"). Important messages to the group into the message file ( e.g. BK\$LOGIN:NOTICE.TXT )

### SYSTEM\_layout

#### Automatic\_backup

Due to a recent system reconfiguration (new system disk and new release of VMS) the automatic backup utility is not available. It is recommended that users do their own backups in the same way as the automatic backup did before by including

a first-level subdirectory such as

[meissb.backup...]\*.\*

on the private directories. It is recommended that users copy important (and only important !) data to this subdirectory for daily full and/or incremental backups.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

BKHELP TELEPHONE (14 July 1986):

Telephone

BIG KARL PHONE DIRECTORY

\*\*\*\*\*

TELEPHONE

BODE

CHR.BODE            5615 -- PRIV: 02461/1250

TELEPHONE

SEHL

G. SEHL            5601 -- PRIV: 0228/311975

TELEPHONE

v.ROSSEN

P.v.Rossen            3096 -- priv: 0228/210900

TELEPHONE

BERG

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

G.P.A.BERG 4121 -- PRIV: 0221/464041

TELEPHONE

BRINKMOELLER

B.BRINKMOELLER 5884 -- priv: 055703

TELEPHONE

HADAMEK

H.HADAMEK 6352 -- PRIV: 02517

TELEPHONE

HARDT

A.HARDT 3623 -- PRIV: 052140

TELEPHONE

HLAWATSCH

G.HLAWATSCH 5601 -- PRIV: 02421/15486

1-86-2569 KFA Direktwahl

1-86-2583

Technische Universitaet Muenchen, Dep. E-18

TELEPHONE

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

KOEHLER

M.KOEHLER 3283 -- PRIV: 053709

TELEPHONE

KRUCK

K.P.KRUCK 6593 -- PRIV: 02402/20980

TELEPHONE

MARTIN

S.MARTIN 3623 -- PRIV: 08544

TELEPHONE

MEISSBURGER

J.MEISSBURGER 4739/5139 -- PRIV: 04853

TELEPHONE

PAUL

D.PAUL 5139 -- PRIV: 056611

TELEPHONE

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

PFEIFFER

J.PFEIFFER      6980 -- PRIV: 02421/53417

TELEPHONE

PRASUHN

D.PRASUHN      4121 -- PRIV: 02244/3682

TELEPHONE

RETZ

A.RETZ      6360 -- PRIV: 054528

TELEPHONE

ROGGE

M.ROGGE      4122 -- PRIV: 08358

TELEPHONE

ROSSKAMP

J.ROSSKAMP      4267 -- PRIV: 08630

TELEPHONE

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

STOFF

H.STOFF                      6490 -- PRIV: 02464/7841

TELEPHONE

EXPERIMENT

BIG KARL CONTROL ROOM (ROOM 47)                      5139  
BIG KARL NORTH HALL (SPECTROMETER)              6374

TELEPHONE

CYCLOTRON

CYCLOTRON CONTROL ROOM (OPERATORS)              6454

TELEPHONE

DEC

DEC-TECHNIKER      4093  
DEC KOELN              0221-5486-1  
DEC Muenchen (RDC center)              0-089-418016

TELEPHONE

TSS

TSS CONSOLE              4380

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

TELEPHONE

KLIMAAANLAGE

KLIMAAANLAGE 6111

TELEPHONE

STRAHLENSCHUTZ

STRAHLENSCH. 6354 -- PRIV: (PROBST) 02428/1472

TELEPHONE

TEKTRONIX

TEKTRONIX 0221/77221

TELEPHONE

SILENA

SILENA ADC's 06055-4021

BKHELP TERMINALS (14 July 1986):

## Terminals

A number of different terminal types is supported around the Big Karl VAX. Special configuration support is available for most of them in a user-transparent way on either DCL command level, Fortran program level or in the form of logical files which may be included in user-generated data files. Some of the more frequently used libraries such as "GMA10" include this support automatically.

## Big Karl terminal type definitions:

/FT1	=	GMA102 - DMA graphics terminal (dev_class = realtime)
/FT2	=	Hewlett Packard 2648A graphics terminal
/FT3	=	Tektronix 4010/4012/4014 graphics terminal
/FT4	=	Westward 1015 graphics terminal
/FT5	=	Hewlett Packard 2623A ANSI/graphics terminal
/FT6	=	Tektronix 4105 color graphics terminal
/FT7	=	Plessey PT-100G high resolution graphics terminal
/VT100	=	VT100 alpha or Retrographics VT100 GB graphics terminal
/LA100	=	DEC LA-100 letter matrix printer (memo-quality)
/LN03	=	DEC LN03 high quality laser printer

To get a list of the terminals installed in the system see:

```
$ TERMINALS
```

TERMINALS

TTYTYPE



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

There exists (BK)-system support to require the terminal type of the log-on terminal SYS\$COMMAND or any other terminal (or logic device) in the system:

1) At DCL command level:

```
$ TTTYTYPE [logical_name]
```

This defines the two symbols:

TT_TYPE	= actual setting of device type (e.g. "VT100")
TT_SYS_TYPE	= true terminal device type (e.g. "FT6" or "DSPORT" for dataswitch lines)

In addition the system defines the symbol:

TT_SAV_TYPE	= permanent setting of SYS\$COMMAND device type at LOGIN time (not altered by calling TTTYTYPE !)
-------------	--

Special a) Any terminal of unknown type will return "TTYUNKN"

Special b) The GMA102 DMA display is returned as TT\_TYPE = "GMAO"

Special c) If a required device is n o t a terminal the device class will be returned instead (e.g. \$ TTTYTYPE USERDISK: would return TT\_TYPE = "DISK").

Special d) Devices not known to the system are returned "UNKNOWN"

Special e) Calling TTTYTYPE in a batch job will return TT\_TYPE = "BATCH"

2) At Fortran program level:

```
CALL BK_TT_TYPE (....)
```

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

For details see:       \$ BKH BKLIB BKUTY BK\_TT\_TYPE

## TERMINALS

### TTMODE

The (BK)-system supports device-independent terminal configuration.

- 1)       At DCL command level:

```
$ TTMODE mode_identifier [logical_name]
```

where "mode\_identifier" is a mode identifier and "logical\_name" is a logical name of a free (\$ SH TERM logical\_name should have "NO OWNER") terminal. If logical\_name is omitted the default "SYS\$COMMAND" is assumed.

If the required mode is not available on the specified device the command is simply ignored (no error message !). The two most important mode\_identifiers are:

```
$ TTMODE dec_mode       switch the terminal to  
                          DEC-Edit mode
```

and

```
$ TTMODE tcs_mode       to switch the terminal to  
                          TCS compatibility mode
```

- 2)       At Fortran program level:

```
CALL BK_TT_MODE ( .... )
```

For details see: \$ BKHelp BKLIB BKUTY BK\_TT\_MODE

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

## TERMINALS

### TTCONFIG

The (BK)-system supports device-dependent terminal configuration for devices of known type:

- 1) At DCL command level:

```
$ TTCONFIG config_file[,config_file] [logical_name]
```

Example:

```
$ TTCONFIG ft5$clr_alph_displ,ft5$aids_off tta4:
```

would clear the alpha-memory of terminal TTA4: (if TTA4: is of type "FT5") and switch off the "Aids"-menue. TTCONFIG does n o t check for correct terminal type (use \$ TTTYPE if necessary).

Note: TTCONFIG should only be used if special features of a known terminal type are to be supported and if the more general TTMODE procedure does not support the required configuration !

- 2) At Fortran program level:

```
CALL BK_TT_CONFIG ( .... )
```

For more details see: \$BKHelp BKLIB BKUTY BK\_TT\_CONFIG

## TERMINALS

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

LA100

Special support is available for the LA100 letter printer  
spooled to queue SYS\$LETTER. Use:

\$ SHOW LOGICAL LETTER\$\*

to display the corresponding definitions. The most important one is:

letter\$config

which configures the letter printer for letter gothic 12, german letter  
size printing (use with german character set on terminal)

See also: \$ BKHelp RUNOFF

TERMINALS

LN03

Special support is available for the LN03 laser printer, see:

\$ BKHelp printers

# BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

## BKHELP UTILITIES (14 July 1986):

### Utilities

The utility programs to support Big Karl experiments all run on the Big Karl group-accounts BK\_name ,e.g. BK\_ONLINE for on-line data taking. To invoke a program just type the program name at DCL (\$) command level. Most programs accept a HELP command at program level to provide additional information on the program. Normal program exit is ...> EXIT or <CNTRL-Y> in case of serious troubles.

Please watch the display of default directory or use the command \$ SH DEF to keep track of your actual working directory !

### UTILITIES

#### CYCLE

CYCLE is Big Karls RSX-11M control utility running on a PDP-11/40. It is used to set and monitor Big Karl status and keeps a background log of all activities.

Related commands are:

>	or	MCR> RUN CYCLE	to invoke the program
		MCR> STATUS	to display the sepcrometer status
		MCR> CYSTOP	to stop ongoing CYCLE activities

### UTILITIES

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

ACQUIRE

\$ ACON            -->    On-line ACQUIRE program system  
\$ ACOFF           -->    Off-line ACQUIRE

Program system to do data acquisition using Big Karls  
MEMPHIS data acquisition system. For more details see:

\$ BKHelp ACQUIRE...            and  
\$ BKHelp MEMPHIS...

UTILITIES

ACPLOT

\$ ACPlot           -->    Spectrum plotting utility

The program allows plotting of spectra generated by  
ACQUIRE or PLSORT on either the CALCOMP plotter or the systems  
lineprinter/plotter LPAO. Plotting format is variable.

See: \$ BKHelp ACPLOT...

UTILITIES

TURTLE

\$ TURTle           -->    CERN turtle decay

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

Ionoptics program to calculate Monte Carlo generated distributions of particle rays through beam transport systems.

UTILITIES

IGM

\$ IGM      -->      Big Karl Interactive Graphics Monitor

This program system allows the generation of graphical representations of very generally formatted data. It provides batch and hardcopy support and is callable as subsystem from FORTRAN program level.

See: \$ BKHelp IGM...

UTILITIES

MOCAL

\$ MOCa1    -->    Big Karl MORRIS chamber calibration utility.

MOCAL is an interactive program system to generate the data tables for calibration of the Morris MWDC. It works together with ACPARAM, the user parameter definition utility for the on-line process ADON (ACQUIRE and ACQUIRERT).

See: \$ BKHelp MOCAL...

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

UTILITIES

PLSORT

\$ PLSort --> Off-line sorting utility

PLSORT is the general purpose off-line sorting utility that provides transparently organized access to ACQUIRE and MOCAL data tables and software modules.

UTILITIES

PLSTATUS

\$ PLStatus --> Show actual status of private PLSORT system

UTILITIES

BKDAP

\$ BKDap --> Big Karl data analyzing programs

This system of programs is used to further process the spectra generated by ACSPEC in order to generate angular distributions, energy calibrations, theoretical DWUCK calculations etc. The modules are:

\$ ANGULAR

\$ CALIBRATE



BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

\$ CROSSECT  
\$ INPUT  
\$ PLOT  
\$ POTENTIAL  
\$ DWUCK  
\$ INFO

UTILITIES

MATRIX

\$ MATrix --> Big Karl optimization utility "MATRIX"

MATRIX calculates the elements of the BIG KARL TRANSPORT matrix using experimental raytracing. The "aberrations" are determined, optimal multipole settings and line broadening are calculated.

UTILITIES

MAGSET

\$ MAGset --> Big Karl magnet setup utility "MAGSET"

MAGSET uses TRANSPORT calculations and empirical data obtained in previous optimization runs to give approximate start values for magnetic field settings. K-value tuning is taken into account and appropriate values are calculated for Q2 and QB.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

UTILITIES

TRANSPORT

\$ TRANsport --> Big Karl TRANSPORT program

TRANSPORT version with extended plotting facilities.  
For details ask S.Martin, ASI.

UTILITIES

TAPECOPY

Tape copy utility to read aritrarely formatted tapes into  
Files-11 disk files (e.g. IBM, HP tapes etc.)

UTILITIES

REVMOC

\$ REVmoc --> Ionoptics code "REVMOC"

UTILITIES

STRAGGLE

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

\$ STRaggle --> Straggling calculation (Vavilov distrib.)

Energy straggling and energy loss simulation.

UTILITIES

BRHO

\$ BRHO --> Relativistic kinematics including energy loss

The program system BRHO is Big Karl's general purpose kinematics utility. It calculates kinematical quantities for two-body reactions including energy loss in composite targets. Output is either numerical or graphical representation of angular and energy dependence. For more details see:

\$ BKHelp BRHO

UTILITIES

KINE

Replaced by BRHO, see: \$ BKHelp BRHO...

UTILITIES

KIN1

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

\$ KIN1       -->       K-values and Q2-tuning

KIN1 calculates the PRE2 (=Q2) values for a matched tuning of Q2 depending on K-value (Lab. angle)

UTILITIES

BKARLO

\$ BKArlo       -->       Energy loss through Big Karl (or  
any other layered detector setup)

BKARLO is a general purpose program to calculate energy loss by integration through composite absorptive layers (such as Big Karl spectrometer or detector setups) It is used for setting up particle identification by energy loss gas counters or plastic scintillators. Input is symbolic with predefined material constant data tables.

See:       \$ BKHelp BKARLO

UTILITIES

MWPC

\$ MWPC       -->       Calculate MWPC voltages

Program to calculate the setup voltages for the Koehler MWPC for standard gas.

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

UTILITIES

GAMES

\$ GAMES --> Big Karl recreation area

Choice of computer games. Warning: When playing one of these games you automatically lower your process priority !

UTILITIES

DEMO

\$ DEMO --> Graphics Demo (TCS-compatible TERMINALS)

Set of demonstration programs to test the TCS (PLOT10) compatibility of a terminal as well as some special enhancements for 41xx-family terminals. For details ask J.Meissburger.

UTILITIES

BKMERGE

Same as BKFIT, see: \$ BKHelp BKFIT...

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

UTILITIES

BKFIT

\$ BKFIT --> Big Karl general purpose spectrum manipulation  
utility to

read , write, merge , modify , analyze and display

spectra written on data files in Big Karl standard format (\*.SIN)

For more details see: \$ BKHelp BKFIT...

Note: BKFIT was planned to become Big Karls most general fitting utility  
by connecting an interactive version of CERN MINUIT with standard and user  
defined fit routines to the user interface provided in BKFIT (J.Meissburger).

UTILITIES

CTR

\$ CTR --> COSYTRIEVE (CTR) data storage and retrieval system  
(Poor man's Datatrieve)

Used to keep literature index or hardware module data bases.

See: \$ BKHelp CTR...

UTILITIES

BIB KARL magnet spectrometer operating system HELP LIBRARY

Author: J.Meissburger

MINUIT

\$ MINUIT --> General purpose fitting utility (CERN)  
with standard extension for Chisquare  
fitting and display of results (MINFIT)

